

SEQUENCE LISTING

<110> Li, Li Furtak, Kazarzyna Perna, Amanda Patturajan, Meera Shimkets, Richard A Guo, Xiaojia Sasha Casman, Stacie J Burgess, Catherine E Malyankar, Uriel M Tchernev, Velizar T Vernet, Corrine A Spytek, Kimberly A Agee, Michele Rastelli, Luca Shenoy, Suresh G Grosse, William M Alsobrook II, John P Lepley, Denise M Gerlach, Valerie Edinger, Schlomit MacDougall, John R Peyman, John A Gunther, Erik Stone, David J Ellerman, Karen Gangolli, Esha A

- <120> Novel Human Proteins, Polynucleotides Encoding Them and Methods of Using the Same
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Ser Arg Ser Leu Pro Arg Val Ser His Val Cys Ser Gln Thr Asn Gly 580 585 590

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Pro His Leu Asp Gly Ala Gly Gly Gly Ala Ala Gly Glu Asp Gly Asp 740 745 750

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Glu Leu Arg Leu Thr Asp Asn Phe Ile Ala Ala Val Arg Arg Asp 100 105 110

Phe Ala Asn Met Thr Ser Leu Val His Leu Thr Leu Ser Arg Asn Thr 115 120 125

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Thr Lys Ala Arg Pro Gln Lys Tyr Glu Gln Leu Leu His Ile Glu Asp
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Asn Asp Phe Ala Met Arg Pro Gly Phe Gly Gly Ser Pro Val Pro Val 65 70 75 80

Gly Ile Asp Val His Val Glu Ser Ile Asp Ser Ile Ser Glu Thr Asn 85 90 95

Met Asp Phe Thr Met Thr Phe Tyr Leu Arg His Tyr Trp Lys Asp Glu 100 105 110

Arg Leu Ser Phe Pro Ser Thr Ala Asn Lys Ser Met Thr Phe Asp His

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His 225	Met	Ser	Leu	Ser	Gln 230	Phe	Phe	Ile	Glu	Asp 235	Phe	Ser	Ala	Ser	Ser 240
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Phe Asp Gly Cys Tyr His Asp Ser Glu Ile Asp Met Asp Gln Thr Ser

Leu Ser Leu Asn Ser Glu Asp Phe Met Arg Arg Lys Ser Ile Cys Ser 385 390 395 400

Pro Ser Thr Asp Ser Ser Arg Ile Lys Arg Arg Lys Ser Leu Gly Gly 405 410 415

His Val Gly Arg Ile Ile Leu Glu Asn Asn His Val Ile Asp Thr Tyr 420 425 430

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His Gln Pro Ser Ala Leu Gly Tyr Ser Pro Ser Leu His Ile Leu Ala 35 40 45

Ile Gly Thr Arg Ser Gly Ala Ile Lys Leu Tyr Gly Ala Pro Gly Val 50 55 60

Glu Phe Met Gly Leu His Gln Glu Asn Asn Ala Val Thr Gln Ile His Leu Leu Pro Gly Gln Cys Gln Leu Val Thr Leu Leu Asp Asp Asn Ser Leu His Leu Trp Ser Leu Lys Val Lys Gly Gly Ala Ser Glu Leu Gln Glu Asp Glu Ser Phe Thr Leu Arg Gly Pro Pro Gly Ala Ala Pro Ser Ala Thr Gln Ile Thr Val Val Leu Pro His Ser Ser Cys Glu Leu Leu Tyr Leu Gly Thr Glu Ser Gly Asn Val Phe Val Val Gln Leu Pro Ala Phe Arg Ala Leu Glu Asp Arg Thr Ile Ser Ser Asp Ala Val Leu Gln Arg Leu Pro Glu Glu Ala Arg His Arg Arg Val Phe Glu Met Val Glu Ala Leu Gln Glu His Pro Arg Asp Pro Asn Gln Ile Leu Ile Gly Tyr Ser Arg Gly Leu Val Val Ile Trp Asp Leu Gln Gly Ser Arg Val Leu Tyr His Phe Leu Ser Ser Gln Gln Leu Glu Asn Ile Trp Trp Gln Arg Asp Gly Arg Leu Leu Val Ser Cys His Ser Asp Gly Ser Tyr Cys Gln Trp Pro Val Ser Ser Glu Ala Gln Gln Pro Glu Pro Leu Arg Ser Leu Val Pro Tyr Gly Pro Phe Pro Cys Lys Ala Ile Thr Arg Ile Leu Trp Leu Thr Thr Arg Gln Gly Leu Pro Phe Thr Ile Phe Gln Gly Gly Met Pro Arg Ala Ser Tyr Gly Asp Arg His Cys Ile Ser Val Ile His Asp Gly Gln Gln Thr Ala Phe Asp Phe Thr Ser Arg Val Ile Gly Thr Val Arg Phe Trp Asp Ala Ser Gly Val Cys Leu Arg Leu Leu Tyr Lys Leu Ser Thr Val Arg Val Phe Leu Thr Asp Thr Asp Pro Asn Glu Asn Phe Ser Ala Gln Gly Glu Asp Glu Trp Pro Pro Leu Arg Lys Val Gly Ser Phe Asp Pro Tyr Ser Asp Asp Pro Arg Leu Gly Ile Gln Lys Ile Phe Leu Cys Lys Tyr Ser Gly Tyr Leu Ala Val Ala Gly Thr Ala Gly Gln Val Leu Val Leu Glu Leu Asn Asp Glu Ala Ala Glu Gln Ala Val Glu Gln Val Glu Ala Asp Leu Leu Gln Asp Gln Glu Gly Tyr Arg Trp Lys Gly His Glu Arg Leu Ala Ala Arg Ser Gly Pro Val Arg Phe Glu Pro Gly Phe Gln Pro Phe Val Leu Val Gln Cys Gln Pro Pro Ala Val Val Thr Ser Leu Ala Leu His Ser Glu Trp Arg Leu Val Ala Phe Gly Thr ,485 Ser His Gly Phe Gly Leu Phe Asp His Gln Gln Arg Arg Gln Val Phe Val Lys Cys Thr Leu His Pro Ser Asp Gln Leu Ala Leu Glu Gly Pro Leu Ser Arg Val Lys Ser Leu Lys Lys Ser Leu Arg Gln Ser Phe Arg Arg Met Arg Arg Ser Arg Val Ser Ser Arg Lys Arg His Pro Ala Gly Pro Pro Gly Glu Ala Gln Glu Gly Ser Ala Lys Ala Glu Arg Pro Gly

Leu Gln Asn Met Glu Leu Ala Pro Val Gln Arg Lys Ile Glu Ala Arg Ser Ala Glu Asp Ser Phe Thr Gly Phe Val Arg Thr Leu Tyr Phe Ala Asp Thr Tyr Leu Lys Asp Ser Ser Arg His Cys Pro Ser Leu Trp Ala Gly Thr Asn Gly Gly Thr Ile Tyr Ala Phe Ser Leu Arg Val Pro Pro Ala Glu Arg Arg Met Asp Glu Pro Val Arg Ala Glu Gln Ala Lys Glu Ile Gln Leu Met His Arg Ala Pro Val Val Gly Ile Leu Val Leu Asp Gly His Ser Val Pro Leu Pro Glu Pro Leu Glu Val Ala His Asp Leu Ser Lys Ser Pro Asp Met Gln Gly Ser His Gln Leu Leu Val Val Ser Glu Glu Gln Phe Lys Val Phe Thr Leu Pro Lys Val Ser Ala Lys Leu Lys Leu Lys Leu Thr Ala Leu Glu Gly Ser Arg Val Arg Arg Val Ser Val Ala His Phe Gly Ser Arg Arg Ala Glu Asp Tyr Gly Glu His His Leu Ala Val Leu Thr Asn Leu Gly Asp Ile Gln Val Val Ser Leu Pro Leu Leu Lys Pro Gln Val Arg Tyr Ser Cys Ile Arg Arg Glu Asp Val Ser Gly Ile Ala Ser Cys Val Phe Thr Lys Tyr Gly Gln Gly Phe Tyr Leu Ile Ser Pro Ser Glu Phe Glu Arg Phe Ser Leu Ser Thr Lys Trp Leu Val Glu Pro Arg Cys Leu Val Asp Ser Ala Glu Thr Lys Asn His

Arg Pro Gly Asn Gly Ala Gly Pro Lys Lys Ala Pro Ser Arg Ala Arg 835 840 845

Asn Ser Gly Thr Gln Ser Asp Gly Glu Glu Lys Gln Pro Gly Leu Val 850 855 860

Met Glu Arg Ala Leu Leu Ser Asp Glu Arg Ala Ala Thr Gly Val His 865 870 875 880

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<211> 3492

<212> DNA

<213> Homo sapiens

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His Gln Pro Ser Ala Leu Gly Tyr Ser Pro Ser Leu His Ile Leu Ala

Val Pro Tyr Gly Pro Phe Pro Cys Lys Ala Ile Thr Arg Ile Leu Trp

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Glu His His Leu Ala Val Leu Thr Asn Leu Gly Asp Ile Gln Val Val 865 870 875 880

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Lys Asn His Arg Pro Gly Asn Gly Ala Gly Pro Lys Lys Ala Pro Ser 945 950 955 960

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Tyr Leu Glu Thr Gly Arg Gly Leu Cys Phe Pro Leu Val Lys Ala Cys 65 70 75 80

Val Phe Gly Asn Glu Pro Lys Ala Ser Asp Glu Val Pro Leu Ala Pro 85 90 95

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Gly Thr Gly Val Val Pro Ala Ala Leu Gly Gly Ala Gly Val Met Ala 340 345 350 Leu Leu Cys Ile Cys Leu Cys Leu Ile Phe Phe Leu Ile Val Lys Ala 355 360 365 Arg Arg Lys Gln Ala Ala Gly Arg Pro Glu Lys Met Asp Asp Glu Asp 370 375 380 Pro Ile Met Gly Thr Ile Thr Ser Gly Ser Arg Lys Lys Pro Trp Pro 385 390 395 Asp Ser Pro Gly Asp Gln Ala Ser Pro Pro Gly Asp Ala Pro Pro Leu 405 410 Glu Glu Gln Lys Glu Leu His Tyr Ala Ser Leu Ser Phe Ser Glu Met 425 Lys Ser Arg Glu Pro Lys Asp Gln Glu Ala Pro Ser Thr Thr Glu Tyr 440 445 Ser Glu Ile Lys Thr Ser Lys 450 455 <210> 30 <211> 1811 <212> DNA <213> Homo sapiens <400> 30 acaaateett etgttgaact etactgtgte aggecageet gagtteattt eteettgage 60 aggaacagtt catggacgaa ctctgaggac cattctgagg acaagaggca tccagtgtca 120 tgagtggaac atgcagcatt ttatggctac agagttaagg caagggttga attccacgag 180 tcaaaaagca gcccttttca gagacccaac tctctggggt gctcaggggc ttgggctgga 240 ttgagaagaa aactgacaag agtaagctgc cctctcttct ctggccatct cacaaaccac 300 agtgcgggcc aactggtcct gcctctttac cacacagaac caagcactag ggataagaca 360 gctgcccatg gtgtccgcgg cgggtctctc tggggatggc aagatgcgag gggtgctcct 420 ggtgctgctc ggccttctct attcttccac cagttgtggc gtccagaaag cttccgtttt 480 ctacggtcct gaccccaagg agggcttggt cagcagcatg gagttcccgt gggtggtgtc 540 gctgcaggac tcccagtaca cacacctggc tttcggctgc atcctgagcg agttctgggt 600 cctcagcatc gcatccgcca ttcagaacag gaaggacatt gtcgttatag tgggtataag 660 taacatggat cctagcaaga ttgctcacac agagtatcca gtcaatacca tcatcataca 720 tgaggacttt gataacaact ccatgagcaa caacatagcc ctcctgaaga cagacacagc 780 gatgcatttt ggcaacctgg tccagtccat ctgcttcctc ggcagaatgc tgcatacacc 840

accagtettg cagaactget gggtgteagg atggaateee acatetgeaa caggaaatea 900 catgaegatg agtgteetga ggaaaatett egtgaaagat ettgaeatgt gteecetata 960 caaacteeag aagaeagaat geggeageea caegaaagag gaaaceaaga etgeetgett 1020

<210> 31

<211> 395

<212> PRT

<213> Homo sapiens

<400> 31

Met Val Ser Ala Ala Gly Leu Ser Gly Asp Gly Lys Met Arg Gly Val 1 5 10 15

Leu Leu Val Leu Leu Gly Leu Leu Tyr Ser Ser Thr Ser Cys Gly Val
20 25 30

Gln Lys Ala Ser Val Phe Tyr Gly Pro Asp Pro Lys Glu Gly Leu Val 35 40 45

Ser Ser Met Glu Phe Pro Trp Val Val Ser Leu Gln Asp Ser Gln Tyr 50 55 60

Thr His Leu Ala Phe Gly Cys Ile Leu Ser Glu Phe Trp Val Leu Ser 65 70 75 80

Ile Ala Ser Ala Ile Gln Asn Arg Lys Asp Ile Val Val Ile Val Gly
85 90 95

Ile Ser Asn Met Asp Pro Ser Lys Ile Ala His Thr Glu Tyr Pro Val
100 105 110

Asn Thr Ile Ile Ile His Glu Asp Phe Asp Asn Asn Ser Met Ser Asn 115 120 125

Asn Ile Ala Leu Leu Lys Thr Asp Thr Ala Met His Phe Gly Asn Leu 130 135 140

Val 145	Gln	Ser	Ile	Cys	Phe 150	Leu	Gly	Arg	Met	Leu 155	His	Thr	Pro	Pro	Val 160
Leu	Gln	Asn	Cys	Trp 165	Val	Ser	Gly	Trp	Asn 170	Pro	Thr	Ser	Ala	Thr 175	Gly
Asn	His	Met	Thr 180	Met	Ser	Val	Leu	Arg 185	Lys	Ile	Phe	Val	Lys 190	Asp	Leu
Asp	Met	Cys 195	Pro	Leu	Tyr	Lys	Leu 200	Gln	Lys	Thr	Glu	Cys 205	Gly	Ser	His
Thr	Lys 210	Glu	Glu	Thr	Lys	Thr 215	Ala	Cys	Leu	Gly	Asp 220	Pro	Gly	Ser	Pro
Met 225	Met	Cys	Gln	Leu	Gln 230	Gln	Phe	Asp	Leu	Trp 235	Val	Leu	Arg	Gly	Ile 240
Leu	Asn	Phe	Gly	Gly 245	Glu	Thr	Cys	Pro	Gly 250	Leu	Phe	Leu	Tyr	Thr 255	Lys
Val	Glu	Asp	Tyr 260	Ser	Lys	Trp	Ile	Thr 265	Ser	Lys	Ala	Glu	Arg 270	Ala	Gly
Pro	Pro	Leu 275	Ser	Ser	Leu	His	His 280	Trp	Glu	Lys	Leu	Ile 285	Ser	Phe	Ser
His	His 290	Gly	Pro	Asn	Ala	Ala 295	Met	Thr	Gln	Lys	Thr 300	Tyr	Ser	Asp	Ser
Glu 305	Leu	Gly	His	Val	Gly 310	Ser	Tyr	Leu	Gln	Gly 315	Gln	Arg	Arg	Thr	Ile 320
Thr	His	Ser	Arg	Leu 325	Gly	Asn	Ser	Ser	Arg 330	Asp	Ser	Leu	Asp	Val 335	Arg
Glu	Lys	Asp	Val 340	Lys	Glu	Ser	Gly	Arg 345	Ser	Pro	Glu	Ala	Ser 350	Val	Gln
Pro	Leu	Tyr 355	Tyr	Asp	Tyr	Tyr	Gly 360	Gly	Glu	Val	Gly	Glu 365	Gly	Arg	Ile
Phe	Ala 370	Gly	Gln	Asn	Arg	Leu 375	Tyr	Gln	Pro	Glu	Glu 380	Ile	Ile	Leu	Val
Ser 385	Phe	Val	Leu	Val	Phe 390	Phe	Cys	Ser	Ser	Ile 395					

<210> 32

<211> 558

<212> PRT

<213> Mus musculus

<400> 32

Met Ala His Ser Glu Glu Gln Ala Ala Val Pro Cys Ala Phe Ile Arg 1 5 10 15

Gln Asn Ser Gly Asn Ser Ile Ser Leu Asp Phe Glu Pro Asp Thr Glu 20 25 30

Tyr Gln Phe Val Glu Gln Leu Glu Glu Arg Tyr Lys Cys Ala Phe Cys 35 40 45

His Ser Val Leu His Asn Pro His Gln Thr Gly Cys Gly His Arg Phe 50 55 60

Cys Gln Gln Cys Ile Arg Ser Leu Arg Glu Leu Asn Ser Val Pro Ile 65 70 75 80

Cys Pro Val Asp Lys Glu Val Ile Lys Pro Gln Glu Val Phe Lys Asp 85 90 95

Asn Cys Cys Lys Arg Glu Val Leu Asn Leu His Val Tyr Cys Lys Asn 100 105 110

Ala Pro Gly Cys Asn Ala Arg Ile Ile Leu Gly Arg Phe Gln Asp His 115 120 125

Leu Gln His Cys Ser Phe Gln Ala Val Pro Cys Pro Asn Glu Ser Cys 130 135 140

Arg Glu Ala Met Leu Arg Lys Asp Val Lys Glu His Leu Ser Ala Tyr 145 150 155 160

Cys Arg Phe Arg Glu Glu Lys Cys Leu Tyr Cys Lys Arg Asp Ile Val 165 170 175

Val Thr Asn Leu Gln Asp His Glu Glu Asn Ser Cys Pro Ala Tyr Pro 180 185 190

Val Ser Cys Pro Asn Arg Cys Val Gln Thr Ile Pro Arg Ala Arg Val 195 200 205

Asn Glu His Leu Thr Val Cys Pro Glu Ala Glu Gln Asp Cys Pro Phe

Lys 225	His	Tyr	Gly	Cys	Thr 230	Val	Lys	Gly	Lys	Arg 235	Gly	Asn	Leu	Leu	Glu 240
His	Glu	Arg	Ala	Ala 245	Leu	Gln	Asp	His	Met 250	Leu	Leu	Val	Leu	Glu 255	Lys
Asn	Tyr	Gln	Leu 260	Glu	Gln	Arg	Ile	Ser 265	Asp	Leu	Tyr	Gln	Ser 270	Leu	Glu
Gln	Lys	Glu 275	Ser	Lys	Ile	Gln	Gln 280	Leu	Ala	Glu	Thr	Val 285	Lys	Lys	Phe
Glu	Lys 290	Glu	Leu	Lys	Gln	Phe 295	Thr	Gln	Met	Phe	Gly 300	Arg	Asn	Gly	Thr
Phe 305	Leu	Ser	Asn	Val	Gln 310	Ala	Leu	Thr	Ser	His 315	Thr	Asp	Lys	Ser	Ala 320
Trp	Leu	Glu	Ala	Gln 325	Val	Arg	Gln	Leu	Leu 330	Gln	Ile	Val	Asn	Gln 335	Gln
Pro	Ser	Arg	Leu 340	Asp	Leu	Arg	Ser	Leu 345	Val	Asp	Ala	Val	Asp 350	Ser	Val
Lys	Gln	Arg 355	Ile	Thr	Gln	Leu	Glu 360	Ala	Ser	Asp	Gln	Arg 365	Leu	Val	Leu
Leu	Gl.u 370	Gly	Glu	Thr	Ser	Lys 375	His	Asp	Ala	His	Ile 380	Asn	Ile	His	Lys
Ala 385	Gln	Leu	Asn	Lys	Asn 390	Glu	Glu	Arg	Phe	Lys 395	Gln	Leu	Glu	Gly	Ala 400
Cys	Tyr	Ser	Gly	Lys 405	Leu	Ile	Trp	Lys	Val 410	Thr	Asp	Tyr	Arg	Val 415	Lys
Lys	Arg	Glu	Ala 420	Val	Glu	Gly	His	Thr 425	Val	Ser	Val	Phe	Ser 430	Gln	Pro
Phe	Tyr	Thr 435	Ser	Arg	Cys	Gly	Tyr 440	Arg	Leu	Cys	Ala	Arg 445	Ala	Tyr	Leu
Asn	Gly 450	Asp	Gly	Ser	Gly	Lys 455	Gly	Thr	His	Leu	Ser 460	Leu	Tyr	Phe	Val

Val Met Arg Gly Glu Phe Asp Ser Leu Leu Gln Trp Pro Phe Arg Gln

Arg Val Thr Leu Met Leu Leu Asp Gln Ser Gly Lys Lys Asn His Ile 485 490 495

Val Glu Thr Phe Lys Ala Asp Pro Asn Ser Ser Phe Lys Arg Pro 500 505 510

Asp Gly Glu Met Asn Ile Ala Ser Gly Cys Pro Arg Phe Val Ser His 515 520 525

Ser Thr Leu Glu Asn Ser Lys Asn Thr Tyr Ile Lys Asp Asp Thr Leu 530 535 540

Phe Leu Lys Val Ala Val Asp Leu Thr Asp Leu Glu Asp Leu 545 555

<210> 33

<211> 558

<212> PRT

<213> Mus musculus

<400> 33

Met Ala His Ser Glu Glu Gln Ala Ala Val Pro Cys Ala Phe Ile Arg
1 5 10 15

Gln Asn Ser Gly Asn Ser Ile Ser Leu Asp Phe Glu Pro Asp Thr Glu 20 25 30

Tyr Gln Phe Val Glu Gln Leu Glu Glu Arg Tyr Lys Cys Ala Phe Cys 35 40 45

His Ser Val Leu His Asn Pro His Gln Thr Gly Cys Gly His Arg Phe 50 55 60

Cys Gln Gln Cys Ile Arg Ser Leu Arg Glu Leu Asn Ser Val Pro Ile 65 70 75 80

Cys Pro Val Asp Lys Glu Val Ile Lys Pro Gln Glu Val Phe Lys Asp 85 90 95

Asn Cys Cys Lys Arg Glu Val Leu Asn Leu His Val Tyr Cys Lys Asn 100 105 110

Ala Pro Gly Cys Asn Ala Arg Ile Ile Leu Gly Arg Phe Gln Asp His 115 120 125

Leu	Gln 130	His	Cys	Ser	Phe	Gln 135	Ala	Val	Pro	Cys	Pro 140	Asn	Glu	Ser	Cys
Arg 145	Glu	Ala	Met	Leu	Arg 150	Lys	Asp	Val	Lys	Glu 155	His	Leu	Ser	Ala	Tyr 160
Cys	Arg	Phe	Arg	Glu 165	Glu	Lys	Cys	Leu	Tyr 170	Cys	Lys	Arg	Asp	Ile 175	Val
Val	Thr	Asn	Leu 180	Gln	Asp	His	Glu	Glu 185	Asn	Ser	Cys	Pro	Ala 190	Tyr	Pro
Val	Ser	Cys 195	Pro	Asn	Arg	Cys	Val 200	Gln	Thr	Ile	Pro	Arg 205	Ala	Arg	Val
Asn	Glu 210	His	Leu	Thr	Val	Cys 215	Pro	Glu	Ala	Glu	Gln 220	Asp	Cys	Pro	Phe
Lys 225	His	Tyr	Gly	Cys	Thr 230	Val	Lys	Gly	Lys	Arg 235	Gly	Asn	Leu	Leu	Glu 240
His	Glu	Arg	Ala	Ala 245	Leu	Gln	Asp	His	Met 250	Leu	Leu	Val	Leu	Glu 255	Lys
Asn	Tyr	Gln	Leu 260	Glu	Gln	Arg	Ile	Ser 265	Asp	Leu	Tyr	Gln	Ser 270	Leu	Glu
Gln	Lys	Glu 275	Ser	Lys	Ile	Gln	Gln 280	Leu	Ala	Glu	Thr	Val 285	Lys	Lys	Phe
Glu	Lys 290	Glu	Leu	Lys	Gln	Phe 295	Thr	Gln	Met	Phe	Gly 300	Arg	Asn	Gly	Thr
Phe 305	Leu	Ser	Asn	Val	Gln 310	Ala	Leu	Thr	Ser	His 315	Thr	Asp	Lys	Ser	Ala 320
Trp	Leu	Glu	Ala	Gln 325	Val	Arg	His	Leu	Leu 330	Gln	Ile	Val	Asn	Gln 335	Gln
Pro	Ser	Arg	Leu 340	Asp	Leu	Arg	Ser	Leu 345	Val	Asp	Ala	Val	Asp 350	Ser	Val
Lys	Gln	Arg 355	Ile	Thr	Gln	Leu	Glu 360	Ala	Ser	Asp	Gln	Arg 365	Leu	Val	Leu
Leu	Glu 370	Gly	Glu	Thr	Ser	Lys 375	His	Asp	Ala	His	Ile 380	Asn	Ile	His	Lys

Ala Gln Leu Asn Lys Asn Glu Glu Arg Phe Lys Gln Leu Glu Gly Ala 385 390 395 400

Cys Tyr Ser Gly Lys Leu Ile Trp Lys Val Thr Asp Tyr Arg Val Lys 405 410 415

Lys Arg Glu Ala Val Glu Gly His Thr Val Ser Val Phe Ser Gln Pro 420 425 430

Phe Tyr Thr Ser Arg Cys Gly Tyr Arg Leu Cys Ala Arg Ala Tyr Leu 435 440 445

Asn Gly Asp Gly Ser Gly Lys Gly Thr His Leu Ser Leu Tyr Phe Val 450 455 460

Val Met Arg Gly Glu Phe Asp Ser Leu Leu Gln Trp Pro Phe Arg Gln 465 470 475 480

Arg Val Thr Leu Met Leu Leu Asp Gln Ser Gly Lys Lys Asn His Ile 485 490 495

Val Glu Thr Phe Lys Ala Asp Pro Asn Ser Ser Phe Lys Arg Pro 500 505 510

Asp Gly Glu Met Asn Ile Ala Ser Gly Cys Pro Arg Phe Val Ser His 515 520 525

Ser Thr Leu Glu Asn Ser Lys Asn Thr Tyr Ile Lys Asp Asp Thr Leu 530 540

Phe Leu Lys Val Ala Val Asp Leu Thr Asp Leu Glu Asp Leu 545 550 555

<210> 34

<211> 557

<212> PRT

<213> Homo sapiens

<400> 34

Met Ala Tyr Ser Glu Glu His Lys Gly Met Pro Cys Gly Phe Ile Arg 1 5 10 15

Gln Asn Ser Gly Asn Ser Ile Ser Leu Asp Phe Glu Pro Ser Ile Glu 20 25 30

Tyr Gln Phe Val Glu Arg Leu Glu Glu Arg Tyr Lys Cys Ala Phe Cys 35 40 45

His Ser Val Leu His Asn Pro His Gln Thr Gly Cys Gly His Arg Phe Cys Gln His Cys Ile Leu Ser Leu Arg Glu Leu Asn Thr Val Pro Ile Cys Pro Val Asp Lys Glu Val Ile Lys Ser Gln Glu Val Phe Lys Asp Asn Cys Cys Lys Arg Glu Val Leu Asn Leu Tyr Val Tyr Cys Ser Asn Ala Pro Gly Cys Asn Ala Lys Val Ile Leu Gly Arg Tyr Gln Asp His Leu Gln Gln Cys Leu Phe Gln Pro Val Gln Cys Ser Asn Glu Lys Cys Arg Glu Pro Val Leu Arg Lys Asp Leu Lys Glu His Leu Ser Ala Ser Cys Gln Phe Arg Lys Glu Lys Cys Leu Tyr Cys Lys Lys Asp Val Val Val Ile Asn Leu Gln Asn His Glu Glu Asn Leu Cys Pro Glu Tyr Pro Val Phe Cys Pro Asn Asn Cys Ala Lys Ile Ile Leu Lys Thr Glu Val Asp Glu His Leu Ala Val Cys Pro Glu Ala Glu Gln Asp Cys Pro Phe Lys His Tyr Gly Cys Ala Val Thr Asp Lys Arg Arg Asn Leu Gln Gln His Glu His Ser Ala Leu Arg Glu His Met Arg Leu Val Leu Glu Lys Asn Val Gln Leu Glu Gln Ile Ser Asp Leu His Lys Ser Leu Glu Gln Lys Glu Ser Lys Ile Gln Gln Leu Ala Glu Thr Ile Lys Lys Leu Glu Lys Glu Phe Lys Gln Phe Ala Gln Leu Phe Gly Lys Asn Gly Ser

Phe 305	Leu	Pro	Asn	Ile	Gln 310	Val	Phe	Ala	Ser	His 315	Ile	Asp	Lys	Ser	Ala 320
Trp	Leu	Glu	Ala	Gln 325	Val	His	Gln	Leu	Leu 330	Gln	Met	Val	Asn	Gln 335	Gln
Gln	Asn	Lys	Phe 340	Asp	Leu	Arg	Pro	Leu 345	Met	Glu	Ala	Val	Asp 350	Thr	Val
Lys	Gln	Lys 355	Ile	Thr	Leu	Leu	Glu 360	Asn	Asn	Asp	Gln	Arg 365	Leu	Ala	Val
Leu	Glu 370	Glu	Glu	Thr	Asn	Lys 375	His	Asp	Thr	His	Ile 380	Asn	Ile	His	Lys
Ala 385	Gln	Leu	Ser	Lys	Asn 390	Glu	Glu	Arg	Phe	Lys 395	Leu	Leu	Glu	Gly	Thr 400
Cys	Tyr	Asn	Gly	Lys 405	Leu	Ile	Trp	Lys	Val 410	Thr	Asp	Tyr	Lys	Met 415	Lys
Lys	Arg	Glu	Ala 420	Val	Asp	Gly	His	Thr 425	Val	Ser	Ile	Phe	Ser 430	Gln	Ser
Phe	Tyr	Thr 435	Ser	Arg	Cys	Gly	Tyr 440	Arg	Leu	Cys	Ala	Arg 445	Ala	Tyr	Leu
Asn	Gly 450	Asp	Gly	Ser	Gly	Arg 455	Gly	Ser	His	Leu	Ser 460	Leu	Tyr	Phe	Val
Val 465	Met	Arg	Gly	Glu	Phe 470	Asp	Ser	Leu	Leu	Gln 475	Trp	Pro	Phe	Arg	Gln 480
Arg	Val	Thr	Leu	Met 485	Leu	Leu	Asp	Gln	Ser 490	Gly	Lys	Lys	Asn	Ile 495	Met
Glu	Thr	Phe	Lys 500	Pro	Asp	Pro	Asn	Ser 505	Ser	Ser	Phe	Lys	Arg 510	Pro	Asp
Gly	Glu	Met 515	Asn	Ile	Ala	Ser	Gly 520	Cys	Pro	Arg	Phe	Val 525	Ala	His	Ser
Val	Leu 530	Glu	Asn	Ala	Lys	Asn 535	Ala	Tyr	Ile	Lys	Asp 540	Asp	Thr	Leu	Phe
Leu 545	Lys	Val	Ala	Val	Asp 550	Leu	Thr	Asp	Leu	Glu 555	Asp	Leu			

<210> 35

<211> 543

<212> PRT

<213> Homo sapiens

<400> 35

Met Glu Ser Ser Lys Lys Met Asp Ser Pro Gly Ala Leu Gln Thr Asn 1 5 10 15

Pro Pro Leu Lys Leu His Thr Asp Arg Ser Ala Gly Thr Pro Val Phe 20 25 30

Val Pro Glu Gln Gly Gly Tyr Lys Glu Lys Phe Val Lys Thr Val Glu 35 40 45

Asp Lys Tyr Lys Cys Glu Lys Cys His Leu Val Leu Cys Ser Pro Lys 50 55 60

Gln Thr Glu Cys Gly His Arg Phe Cys Glu Ser Cys Met Ala Ala Leu 65 70 75 80

Leu Ser Ser Ser Pro Lys Cys Thr Ala Cys Gln Glu Ser Ile Val 85 90 95

Lys Asp Lys Val Phe Lys Asp Asn Cys Cys Lys Arg Glu Ile Leu Ala 100 105 110

Leu Gln Ile Tyr Cys Arg Asn Glu Ser Arg Gly Cys Ala Glu Gln Leu 115 120 125

Thr Leu Gly His Leu Leu Val His Leu Lys Asn Asp Cys His Phe Glu 130 135 140

Glu Leu Pro Cys Val Arg Pro Asp Cys Lys Glu Lys Val Leu Arg Lys 145 150 155 160

Asp Leu Arg Asp His Val Glu Lys Ala Cys Lys Tyr Arg Glu Ala Thr 165 170 175

Cys Ser His Cys Lys Ser Gln Val Pro Met Ile Ala Leu Gln Lys His
180 185 190

Glu Asp Thr Asp Cys Pro Cys Val Val Ser Cys Pro His Lys Cys 195 200 205

Ser Val Gln Thr Leu Leu Arg Ser Glu Gly Thr Asn Gln Gln Ile Lys

Ala 225	His	Glu	Ala	Ser	Ser 230	Ala	Val	Gln	His	Val 235	Asn	Leu	Leu	Lys	Glu 240
Trp	Ser	Asn	Ser	Leu 245	Glu	Lys	Lys	Val	Ser 250	Leu	Leu	Gln	Asn	Glu 255	Ser
Val	Glu	Lys	Asn 260	Lys	Ser	Ile	Gln	Ser 265	Leu	His	Asn	Gln	Ile 270	Cys	Ser
Phe	Glu	Ile 275	Glu	Ile	Glu	Arg	Gln 280	Lys	Glu	Met	Leu	Arg 285	Asn	Asn	Glu
Ser	Lys 290	Ile	Leu	His	Leu	Gln 295	Arg	Val	Ile	Asp	Ser 300	Gln	Ala	Glu	Lys
Leu 305	Lys	Glu	Leu	Asp	Lys 310	Glu	Ile	Arg	Ser	Phe 315	Arg	Gln	Asn	Trp	Glu 320
Glu	Ala	Asp	Ser	Met 325	Lys	Ser	Ser	Val	Glu 330	Ser	Leu	Gln	Asn	Arg 335	Val
Thr	Glu	Leu	Glu 340	Ser	Val	Asp	Lys	Ser 345	Ala	Gly	Gln	Val	Ala 350	Arg	Asn
Thr	Gly	Leu 355	Leu	Glu	Ser	Gln	Leu 360	Ser	Arg	His	Asp	Gln 365	Met	Leu	Ser
Val	His 370	Asp	Ile	Arg	Leu	Ala 375	Asp	Met	Asp	Leu	Arg 380	Phe	Gln	Val	Leu
Glu 385	Thr	Ala	Ser	Tyr	Asn 390	Gly	Val	Leu	Ile	Trp 395	Lys	Ile	Arg	Asp	Tyr 400
Lys	Arg	Arg	Lys	Gln 405	Glu	Ala	Val		Gly 410	Lys	Thr	Leu	Ser	Leu 415	Tyr
Ser	Gln	Pro	Phe 420	Tyr	Thr	Gly	Tyr	Phe 425	Gly	Tyr	Lys	Met	Cys 430	Ala	Arg
Val	Tyr	Leu 435	Asn	Gly	Asp	Gly	Met 440	Gly	Lys	Gly	Thr	His 445	Leu	Ser	Leu
Phe	Phe 450	Val	Ile	Met	Arg	Gly 455	Glu	Tyr	Asp	Ala	Leu 460	Leu	Pro	Trp	Pro
Phe	Lys	Gln	Lys	Val	Thr	Leu	Met	Leu	Met	Asp	Gln	Gly	Ser	Ser	Arg

465 470	475	480
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Arg His Leu Gly Asp Ala Phe Lys Pro Asp Pro Asn Ser Ser Phe 485 490 495

Lys Lys Pro Thr Gly Glu Met Asn Ile Ala Ser Gly Cys Pro Val Phe 500 505 510

Val Ala Gln Thr Val Leu Glu Asn Gly Thr Tyr Ile Lys Asp Asp Thr 515 520 525

Ile Phe Ile Lys Val Ile Val Asp Thr Ser Asp Leu Pro Asp Pro 530 535 540

<210> 36

<211> 568

<212> PRT

<213> Homo sapiens

<400> 36

Met Glu Ser Ser Lys Lys Met Asp Ser Pro Gly Ala Leu Gln Thr Asn 1 5 10 15

Pro Pro Leu Lys Leu His Thr Asp Arg Ser Ala Gly Thr Pro Val Phe 20 25 30

Val Pro Glu Gln Gly Gly Tyr Lys Glu Lys Phe Val Lys Thr Val Glu 35 40 45

Asp Lys Tyr Lys Cys Glu Lys Cys His Leu Val Leu Cys Ser Pro Lys 50 55 60

Gln Thr Glu Cys Gly His Arg Phe Cys Glu Ser Cys Met Ala Ala Leu 65 70 75 80

Leu Ser Ser Ser Pro Lys Cys Thr Ala Cys Gln Glu Ser Ile Val 85 90 95

Lys Asp Lys Val Phe Lys Asp Asn Cys Cys Lys Arg Glu Ile Leu Ala 100 105 110

Leu Gln Ile Tyr Cys Arg Asn Glu Ser Arg Gly Cys Ala Glu Gln Leu 115 120 125

Thr Leu Gly His Leu Leu Val His Leu Lys Asn Asp Cys His Phe Glu 130 135 140

Glu 145	Leu	Pro	Cys	Val	Arg 150	Pro	Asp	Cys	Lys	Glu 155	Lys	Val	Leu	Arg	Lys 160
Asp	Leu	Arg	Asp	His 165	Val	Glu	Lys	Ala	Cys 170	Lys	Tyr	Arg	Glu	Ala 175	Thr
Cys	Ser	His	Cys 180	Lys	Ser	Gln	Val	Pro 185	Met	Ile	Ala	Leu	Gln 190	Lys	His
Glu	Asp	Thr 195	Asp	Cys	Pro	Cys	Val 200	Val	Val	Ser	Cys	Pro 205	His	Lys	Cys
Ser	Val 210	Gln	Thr	Leu	Leu	Arg 215	Ser	Glu	Leu	Ser	Ala 220	His	Leu	Ser	Glu
Cys 225	Val	Asn	Ala	Pro	Ser 230	Thr	Cys	Ser	Phe	Lys 235	Arg	Tyr	Gly	Cys	Val 240
Phe	Gln	Gly	Thr	Asn 245	Gln	Gln	Ile	Lys	Ala 250	His	Glu	Ala	Ser	Ser 255	Ala
Val	Gln	His	Val 260	Asn	Leu	Leu	Lys	Glu 265	Trp	Ser	Asn	Ser	Leu 270	Glu	Lys
Lys	Val	Ser 275	Leu	Leu	Gln	Asn	Glu 280	Ser	Val	Glu	Lys	Asn 285	Lys	Ser	Ile
Gln	Ser 290	Leu	His	Asn	Gln	Ile 295	Cys	Ser	Phe	Glu	Ile 300	Glu	Ile	Glu	Arg
Gln 305	Lys	Glu	Met	Leu	Arg 310	Asn	Asn	Glu	Ser	Lys 315	Ile	Leu	His	Leu	Gln 320
Arg	Val	Ile	Asp	Ser 325	Gln	Ala	Glu	Lys	Leu 330	Lys	Glu	Leu	Asp	Lys 335	Glu
Ile	Arg	Pro	Phe 340	Arg	Gln	Asn	Trp	Glu 345	Glu	Ala	Asp	Ser	Met 350	Lys	Ser
Ser	Val	Glu 355	Ser	Leu	Gln	Asn	Arg 360	Val	Thr	Glu	Leu	Glu 365	Ser	Val	Asp
Lys	Ser 370	Ala	Gly	Gln	Val	Ala 375	Arg	Asn	Thr	Gly	Leu 380	Leu	Glu	Ser	Gln
Leu 385	Ser	Arg	His	Asp	Gln 390	Met	Leu	Ser	Val	His 395	Asp	Ile	Arg	Leu	Ala 400

Asp Met Asp Leu Arg Phe Gln Val Leu Glu Thr Ala Ser Tyr Asn Gly 405 410 415

Val Leu Ile Trp Lys Ile Arg Asp Tyr Lys Arg Arg Lys Gln Glu Ala 420 425 430

Val Met Gly Lys Thr Leu Ser Leu Tyr Ser Gln Pro Phe Tyr Thr Gly 435 440 445

Tyr Phe Gly Tyr Lys Met Cys Ala Arg Val Tyr Leu Asn Gly Asp Gly 450 455 460

Met Gly Lys Gly Thr His Leu Ser Leu Phe Phe Val Ile Met Arg Gly 465 470 475 480

Glu Tyr Asp Ala Leu Leu Pro Trp Pro Phe Lys Gln Lys Val Thr Leu 485 490 495

Met Leu Met Asp Gln Gly Ser Ser Arg Arg His Leu Gly Asp Ala Phe \cdot 500 510

Lys Pro Asp Pro Asn Ser Ser Ser Phe Lys Lys Pro Thr Gly Glu Met 515 520 525

Asn Ile Ala Ser Gly Cys Pro Val Phe Val Ala Gln Thr Val Leu Glu 530 540

Asn Gly Thr Tyr Ile Lys Asp Asp Thr Ile Phe Ile Lys Val Ile Val 545 550 555 560

Asp Thr Ser Asp Leu Pro Asp Pro 565

<210> 37

<211> 159

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: MATH domain sequence

<400> 37

Thr Ile Lys Asn Phe Ser Lys Ile Lys Glu Glu Ala Lys Glu Gly Arg

1 5 10 15

Glu Gly Glu Glu Tyr Tyr Thr Ser Pro Val Glu Glu Arg Phe Asn Ile

25

30

20

Pro Trp Arg Leu Asn Val Leu Arg Ile Tyr Arg Asn Gly Gly Glu
35 40 45

Gly Arg Ser Gly Lys Phe Leu Gly Leu Tyr Leu His Cys Leu Lys Glu
50 55 60

Glu Lys Asp Ser Pro Thr Ile Glu Asn Leu Lys Trp Ser Ile Glu Thr
65 70 75 80

Glu Phe Thr Leu Lys Leu Val Ser Asp Asn Gly Lys Ser Ile Arg Arg 85 90 95

Met Ser Ser Thr Thr Leu Thr Lys Lys Thr Lys Asp Ala Lys Asn Asn 100 105 110

Ser His Val Phe Glu Lys Pro Thr Gly Glu Gly Trp Gly Lys Ser Gly 115 120 125

Phe Lys Lys Phe Ile Ser Trp Asp Asp Leu Glu Asp Asp Tyr Asn Gly 130 135 140

Tyr Leu Val Asp Asp Ser Ile Ile Ile Glu Ala Glu Val Lys Ile 145 150 155

<210> 38

<211> 143

<212> PRT

<213> Homo sapiens

<400> 38

Lys Val Thr Asp Tyr Lys Met Lys Lys Arg Glu Ala Val Asp Gly His

1 5 10 15

Thr Val Ser Ile Phe Ser Gln Ser Phe Tyr Thr Ser Arg Cys Gly Tyr 20 25 30

Arg Leu Cys Ala Arg Ala Tyr Leu Asn Gly Asp Gly Ser Gly Arg Gly
35 40 45

Ser His Leu Ser Leu Tyr Phe Val Val Met Arg Gly Glu Phe Asp Ser 50 55 60

Leu Leu Gln Trp Pro Phe Arg Gln Arg Val Thr Leu Met Leu Leu Asp
65 70 75 80

Gln Ser Gly Lys Lys Asn Ile Met Glu Thr Phe Lys Pro Asp Pro Asn 85 90 95

Ser Ser Ser Phe Lys Arg Pro Asp Gly Glu Met Asn Ile Ala Ser Gly
100 105 110

Cys Pro Arg Phe Val Ala His Ser Val Leu Glu Asn Ala Lys Asn Ala 115 120 125

Tyr Ile Lys Asp Asp Thr Leu Phe Leu Lys Val Ala Val Asp Leu 130 135 140

<210> 39

<211> 700

<212> PRT

<213> Homo sapiens

<400> 39

Leu Thr Asp Asn Phe Ile Ala Ala Val Arg Arg Arg Asp Phe Ala Asn 1 5 10 15

Met Thr Ser Leu Val His Leu Thr Leu Ser Arg Asn Thr Ile Gly Gln
20 25 30

Val Ala Ala Gly Ala Phe Ala Asp Leu Arg Ala Leu Arg Ala Leu His
35 40 45

Leu Asp Ser Asn Arg Leu Ala Glu Val Arg Gly Asp Gln Leu Arg Gly 50 55 60

Leu Gly Asn Leu Arg His Leu Ile Leu Gly Asn Asn Gln Ile Arg Arg
65 70 75 80

Val Glu Ser Ala Ala Phe Asp Ala Phe Leu Ser Thr Val Glu Asp Leu 85 90 95

Asp Leu Ser Tyr Asn Asn Leu Glu Ala Leu Pro Trp Glu Ala Val Gly
100 105 110

Gln Met Val Asn Leu Asn Thr Leu Thr Leu Asp His Asn Leu Ile Asp 115 120 125

His Ile Ala Glu Gly Thr Phe Val Gln Leu His Lys Leu Val Arg Leu 130 135 140

Asp Met Thr Ser Asn Arg Leu His Lys Leu Pro Pro Asp Gly Leu Phe 145 150 155 160

Leu Arg Ser Gln Gly Thr Gly Pro Lys Pro Pro Thr Pro Leu Thr Val Ser Phe Gly Gly Asn Pro Leu His Cys Asn Cys Glu Leu Leu Trp Leu Arg Arg Leu Thr Arg Glu Asp Asp Leu Glu Thr Cys Ala Thr Pro Glu His Leu Thr Asp Arg Tyr Phe Trp Ser Ile Pro Glu Glu Glu Phe Leu Cys Glu Pro Pro Leu Ile Thr Arg Gln Ala Gly Gly Arg Ala Leu Val Val Glu Gly Gln Ala Val Ser Leu Arg Cys Arg Ala Val Gly Asp Pro Glu Pro Val Val His Trp Val Ala Pro Asp Gly Arg Leu Leu Gly Asn Ser Ser Arg Thr Arg Val Arg Gly Asp Gly Thr Leu Asp Val Thr Ile Thr Thr Leu Arg Asp Ser Gly Thr Phe Thr Cys Ile Ala Ser Asn Ala Ala Gly Glu Ala Thr Ala Pro Val Glu Val Cys Val Val Pro Leu Pro Leu Met Ala Pro Pro Pro Ala Ala Pro Pro Pro Leu Thr Glu Pro Gly Ser Ser Asp Ile Ala Thr Pro Gly Arg Pro Gly Ala Asn Asp Ser Ala Ala Glu Arg Arg Leu Val Ala Ala Glu Leu Thr Ser Asn Ser Val Leu Ile Arg Trp Pro Ala Gln Arg Pro Val Pro Gly Ile Arg Met Tyr Gln Val Gln Tyr Asn Ser Ser Val Asp Asp Ser Leu Val Tyr Arg Met Ile Pro Ser Thr Ser Gln Thr Phe Leu Val Asn Asp Leu Ala Ala Gly Arg

Ala Tyr Asp Leu Cys Val Leu Ala Val Tyr Asp Asp Gly Ala Thr Ala Leu Pro Ala Thr Arg Val Val Gly Cys Val Gln Phe Thr Thr Ala Gly Asp Pro Ala Pro Cys Arg Pro Leu Arg Ala His Phe Leu Gly Gly Thr Met Ile Ile Ala Ile Gly Gly Val Ile Val Ala Ser Val Leu Val Phe Ile Val Leu Leu Met Ile Arg Tyr Lys Val Tyr Gly Asp Gly Asp Ser Arg Arg Val Lys Gly Ser Arg Ser Leu Pro Arg Val Ser His Val Cys Ser Gln Thr Asn Gly Ala Gly Thr Gly Ala Ala Gln Ala Pro Ala Leu Pro Ala Gln Asp His Tyr Glu Ala Leu Arg Glu Val Glu Ser Gln Ala Ala Pro Ala Val Ala Val Glu Ala Lys Ala Met Glu Ala Glu Thr Ala Ser Ala Glu Pro Glu Val Val Leu Gly Arg Ser Leu Gly Gly Ser Ala Thr Ser Leu Cys Leu Leu Pro Ser Glu Glu Thr Ser Gly Glu Glu Ser Arg Ala Ala Val Gly Pro Arg Arg Ser Arg Ser Gly Ala Leu Glu Pro Pro Thr Ser Ala Pro Pro Thr Leu Ala Leu Val Pro Gly Gly Ala Ala Ala Arg Pro Arg Pro Gln Gln Arg Tyr Ser Phe Asp Gly Asp Tyr Gly Ala Leu Phe Gln Ser His Ser Tyr Pro Arg Arg Ala Arg Arg Thr Lys Arg His Arg Ser Thr Pro His Leu Asp Gly Ala Gly Gly Ala Ala

Gly Glu Asp Gly Asp Leu Gly Leu Gly Ser Ala Arg Ala Cys Leu Ala 675 680 685

Phe Thr Ser Thr Glu Trp Met Leu Glu Ser Thr Val 690 695 700

<210> 40

<211> 492

<212> PRT

<213> Mus musculus

<400> 40

Met Ala Pro Gly Pro Phe Ser Ser Arg Leu Phe Ser Pro Pro Pro Ala 1 5 10 15

Ala Leu Pro Phe Leu Leu Leu Trp Ala Gly Ala Ser Arg Ser Gln 20 25 30

Pro Cys Pro Gly Arg Cys Ile Cys Gln Asn Val Ala Pro Thr Leu Thr 35 40 45

Met Leu Cys Ala Lys Thr Gly Leu Leu Phe Val Pro Pro Ala Ile Asp 50 55 60

Arg Arg Val Val Glu Leu Arg Leu Thr Asp Asn Phe Ile Ala Ala Val 65 70 75 80

Arg Arg Arg Asp Phe Ala Asn Met Thr Ser Leu Val His Leu Thr Leu 85 90 95

Ser Arg Asn Thr Ile Gly Gln Val Ala Ala Gly Ala Phe Ala Asp Leu 100 105 110

Arg Ala Leu Arg Ala Leu His Leu Asp Ser Asn Arg Leu Ala Glu Val 115 120 125

Arg Gly Asp Gln Leu Arg Gly Leu Gly Asn Leu Arg His Leu Ile Leu 130 135 140

Gly Asn Asn Gln Ile Arg Lys Val Glu Ser Ala Ala Phe Asp Ala Phe 145 150 155 160

Leu Ser Thr Val Glu Asp Leu Asp Leu Ser Tyr Asn Asn Leu Glu Ala 165 170 175

Leu Pro Trp Glu Ala Val Gly Gln Met Val Asn Leu Asn Thr Leu Thr

			180					185					190		
Leu	Asp	His 195	Asn	Leu	Ile	Asp	His 200	Ile	Ala	Glu	Gly	Thr 205	Phe	Val	Gln
Leu	His 210	Lys	Leu	Val	Arg	Leu 215	Asp	Met	Thr	Ser	Asn 220	Arg	Leu	His	Lys
Leu 225	Pro	Pro	Asp	Gly	Leu 230	Phe	Leu	Arg	Ser	Gln 235	Gly	Gly	Gly	Pro	Lys 240
Pro	Pro	Thr	Pro	Leu 245	Thr	Val	Ser	Phe	Gly 250	Gly	Asn	Pro	Leu	His 255	Cys
Asn	Cys	Glu	Leu 260	Leu	Trp	Leu	Arg	Arg 265	Leu	Thr	Arg	Glu	Asp 270	Asp	Leu
Glu	Thr	Cys 275	Ala	Thr	Pro	Glu	His 280	Leu	Thr	Asp	Arg	Tyr 285	Phe	Trp	Ser
Ile	Pro 290	Glu	Glu	Glu	Phe	Leu 295	Cys	Glu	Pro	Pro	Leu 300	Ile	Thr	Arg	Gln
Ala 305	Gly	Gly	Arg	Ala	Leu 310	Val	Val	Glu	Gly	Gln 315	Ala	Val	Ser	Leu	Arg 320
Суз	Arg	Ala	Val	Gly 325	Asp	Pro	Glu	Pro	Val 330	Val	His	Trp	Val	Ala 335	Pro
Asp	Gly	Arg	Leu 340	Leu	Gly	Asn	Ser	Ser 345	Arg	Thr	Arg	Val	Arg 350	Gly	Asp
Gly	Thr	Leu 355	Asp	Val	Thr	Ile	Thr 360	Thr	Leu	Arg	Asp	Ser 365	Gly	Thr	Phe
Thr	Cys 370	Ile	Ala	Ser	Asn	Ala 375	Ala	Gly	Glu	Ala	Thr 380	Ala	Pro	Val	Glu
Val 385	Cys	Val	Val	Pro	Leu 390	Pro	Leu	Met	Ala	Pro 395	Pro	Pro	Ala	Ala	Pro 400
Pro	Pro	Leu	Thr	Glu 405	Pro	Gly	Ser	Ser	Asp 410	Ile	Ala	Thr	Pro	Gly 415	Arg
Pro	Gly	Ala	Asn 420	Asp	Ser	Ala	Thr	Glu 425	Arg	Arg	Leu	Val	Ala 430	Ala	Glu

Leu Thr Ser Ser Ser Val Leu Ile Arg Trp Pro Ala Gln Arg Pro Val

435 440 445

Pro Gly Ile Arg Met Tyr Gln Val Gln Tyr Asn Ser Ser Ala Asp Asp 450 450 460

Ser Leu Val Tyr Ser Ser Ser Cys Pro Gly Thr His Tyr Val Asp Gln 465 470 475 480

Asp Gly Leu Glu Ile Arg Val Pro Leu Ala Ser Ala 485 490

<210> 41

<211> 832

<212> PRT

<213> Homo sapiens

<400> 41

Leu Glu Ser Val Ser Gly Gly Glu Gly Cys Val Ala Glu Pro Gly Ser

1 5 10 15

Pro Gly Ala Pro Arg Ser Arg Pro Arg Cys His Pro Ala Gly Gly Arg
20 25 30

Cys Cys Leu Ala Gln Ala Leu Ser Asp Gln Thr Met Glu Thr Leu Leu 35 40 45

Gly Gly Leu Leu Ala Phe Gly Met Ala Phe Ala Val Val Asp Ala Cys 50 60

Pro Lys Tyr Cys Val Cys Gln Asn Leu Ser Glu Ser Leu Gly Thr Leu 65 70 75 80

Cys Pro Ser Lys Gly Leu Leu Phe Val Pro Pro Asp Ile Asp Arg Arg 85 90 95

Thr Val Glu Leu Arg Leu Gly Gly Asn Phe Ile Ile His Ile Ser Arg 100 105 110

Gln Asp Phe Ala Asn Met Thr Gly Leu Val Asp Leu Thr Leu Ser Arg 115 120 125

Asn Thr Ile Ser His Ile Gln Pro Phe Ser Phe Leu Asp Leu Glu Ser 130 135 140

Leu Arg Ser Leu His Leu Asp Ser Asn Arg Leu Pro Ser Leu Gly Glu
145 150 155 160

Asp	Thr	Leu	Arg	Gly 165	Leu	Val	Asn	Leu	Gln 170	His	Leu	Ile	Val	Asn 175	Asn
Asn	Gln	Leu	Gly 180	Gly	Ile	Ala	Asp	Glu 185	Ala	Phe	Glu	Asp	Phe 190	Leu	Leu
Thr	Leu	Glu 195	Asp	Leu	Asp	Leu	Ser 200	Tyr	Asn	Asn	Leu	His 205	Gly	Leu	Pro
Trp	Asp 210	Ser	Val	Arg	Arg	Met 215	Val	Asn	Leu	His	Gln 220	Leu	Ser	Leu	Asp
His 225	Asn	Leu	Leu	Asp	His 230	Ile	Ala	Glu	Gly	Thr 235	Phe	Ala	Asp	Leu	Gln 240
Lys	Leu	Ala	Arg	Leu 245	Asp	Leu	Thr	Ser	Asn 250	Arg	Leu	Gln	Lys	Leu 255	Pro
Pro	Asp	Pro	Ile 260	Phe	Ala	Arg	Ser	Gln 265	Ala	Ser	Ala	Leu	Thr 270	Ala	Thr
Pro	Phe	Ala 275	Pro	Pro	Leu	Ser	Phe 280	Ser	Phe	Gly	Gly	Asn 285	Pro	Leu	His
Cys	Asn 290	Cys	Glu	Leu	Leu	Trp 295	Leu	Arg	Arg	Leu	Glu 300	Arg	Asp	Asp	Asp
Leu 305	Glu	Thr	Cys	Gly	Ser 310	Pro	Gly	Gly	Leu	Lys 315	Gly	Arg	Tyr	Phe	Trp 320
His	Val	Arg	Glu	Glu 325	Glu	Phe	Val	Cys	Glu 330	Pro	Pro	Leu	Ile	Thr 335	Gln
His	Thr	His	Lys 340	Leu	Leu	Val	Leu	Glu 345	Gly	Gln	Ala	Ala	Thr 350	Leu	Lys
Cys	Lys	Ala 355	Ile	Gly	Asp	Pro	Ser 360	Pro	Leu	Ile	His	Trp 365	Val	Ala	Pro
Asp	Asp 370	Arg	Leu	Val	Gly	Asn 375	Ser	Ser	Arg	Thr	Ala 380	Val	Tyr	Asp	Asn
Gly 385	Thr	Leu	Asp	Ile	Phe 390	Ile	Thr	Thr	Ser	Gln 395	Asp	Ser	Gly	Ala	Phe 400
Thr	Cys	Ile	Ala	Ala 405	Asn	Ala	Ala	Gly	Glu 410	Ala	Thr	Ala	Met	Val 415	Glu

Val Ser Ile Val Gln Leu Pro His Leu Ser Asn Ser Thr Ser Arg Thr Ala Pro Pro Lys Ser Arg Leu Ser Asp Ile Thr Gly Ser Ser Lys Thr Ser Arg Gly Gly Gly Ser Gly Gly Glu Pro Pro Lys Ser Pro Pro Glu Arg Ala Val Leu Val Ser Glu Val Thr Thr Thr Ser Ala Leu Val Lys Trp Ser Val Ser Lys Ser Ala Pro Arg Val Lys Met Tyr Gln Leu Gln Tyr Asn Cys Ser Asp Asp Glu Val Leu Ile Tyr Arg Met Ile Pro Ala Ser Asn Lys Ala Phe Val Val Asn Asn Leu Val Ser Gly Thr Gly Tyr Asp Leu Cys Val Leu Ala Met Trp Asp Asp Thr Ala Thr Thr Leu Thr Ala Thr Asn Ile Val Gly Cys Ala Gln Phe Phe Thr Lys Ala Asp Tyr Pro Gln Cys Gln Ser Met His Ser Gln Ile Leu Gly Gly Thr Met Ile Leu Val Ile Gly Gly Ile Ile Val Ala Thr Leu Leu Val Phe Ile Val Ile Leu Met Val Arg Tyr Lys Val Cys Asn His Glu Ala Pro Ser Lys Met Ala Ala Ala Val Ser Asn Val Tyr Ser Gln Thr Asn Gly Ala Gln Pro Pro Pro Pro Ser Ser Ala Pro Ala Gly Ala Pro Pro Gln Gly Pro Pro Lys Val Val Arg Asn Glu Leu Leu Asp Phe Thr Ala Ser Leu Ala Arg Ala Ser Asp Ser Ser Ser Ser Ser Leu Gly Ser

Gly Glu Ala Ala Gly Leu Gly Arg Ala Pro Trp Arg Ile Pro Pro Ser 675 680 685

Ala Pro Arg Pro Lys Pro Ser Leu Asp Arg Leu Met Gly Ala Phe Ala 690 695 700

Ser Leu Asp Leu Lys Ser Gln Arg Lys Glu Glu Leu Leu Asp Ser Arg 705 710 715 720

Thr Pro Ala Gly Arg Gly Ala Gly Thr Ser Ala Arg Gly His His Ser 725 730 735

Asp Arg Glu Pro Leu Leu Gly Pro Pro Ala Ala Arg Ala Arg Ser Leu 740 745 750

Leu Pro Leu Pro Leu Glu Gly Lys Ala Lys Arg Ser His Ser Phe Asp
755 760 765

Met Gly Asp Phe Ala Ala Ala Ala Gly Gly Val Val Pro Gly Gly 770 775 780

Tyr Ser Pro Pro Arg Lys Val Ser Asn Ile Trp Thr Lys Arg Ser Leu 785 790 795 800

Ser Val Asn Gly Met Leu Leu Pro Phe Glu Glu Ser Asp Leu Val Gly 805 810 815

Ala Arg Gly Thr Phe Gly Ser Ser Glu Trp Val Met Glu Ser Thr Val 820 825 830

<210> 42

<211> 789

<212> PRT

<213> Cynomolgus monkey

<400> 42

Met Glu Thr Leu Leu Gly Gly Leu Leu Ala Phe Gly Met Ala Phe Ala 1 5 10 15

Val Val Asp Ala Cys Pro Lys Tyr Cys Val Cys Gln Asn Leu Ser Glu
20 25 30

Ser Leu Gly Thr Leu Cys Pro Ser Lys Gly Leu Leu Phe Val Pro Pro 35 40 45

Ile His Ile Ser Arg Gln Asp Phe Ala Asn Met Thr Gly Leu Val Asp Leu Thr Leu Ser Arg Asn Thr Ile Ser His Ile Gln Pro Phe Ser Phe Leu Asp Leu Glu Ser Leu Arg Ser Leu His Leu Asp Ser Asn Arg Leu Pro Ser Leu Gly Glu Asp Thr Leu Arg Gly Leu Val Asn Leu Gln His Leu Ile Val Asn Asn Asn Gln Leu Gly Gly Ile Ala Asp Glu Ala Phe Glu Asp Phe Leu Leu Thr Leu Glu Asp Leu Asp Leu Ser Tyr Asn Asn Leu His Gly Leu Pro Trp Asp Ser Val Arg Arg Met Val Asn Leu His Gln Leu Ser Leu Asp His Asn Leu Leu Asp His Ile Ala Glu Gly Thr Phe Ala Asp Leu Gln Lys Leu Ala Arg Leu Asp Leu Thr Ser Asn Arg Leu Gln Lys Leu Pro Pro Asp Pro Ile Phe Ala Arg Ser Gln Ala Ser Ala Leu Thr Ala Thr Pro Phe Ala Pro Pro Leu Ser Phe Ser Phe Gly Gly Asn Pro Leu His Cys Asn Cys Glu Leu Leu Trp Leu Arg Arg Leu Glu Arg Asp Asp Leu Glu Thr Cys Gly Ser Pro Gly Gly Leu Lys Gly Arg Tyr Phe Trp His Val Arg Glu Glu Glu Phe Val Cys Glu Pro Pro Leu Ile Thr Gln His Thr His Lys Leu Val Leu Glu Gly Gln

Asp Ile Asp Arg Arg Thr Val Glu Leu Arg Leu Gly Gly Asn Phe Ile

Ala Ala 305	Thr Leu	Lys Cys		Ala	Ile	Gly	Asp 315	Pro	Ser	Pro	Leu	Ile 320
His Trp	Val Ala	Pro Asp 325	Asp	Arg	Leu	Val 330	Gly	Asn	Ser	Ser	Arg 335	Thr
Ala Val	Tyr Asp 340	Asn Gly	Thr	Leu	Asp 345	Ile	Phe	Ile	Thr	Thr 350	Ser	Gln
Asp Ser	Gly Ala 355	Phe Thr	Cys	Ile 360	Ala	Ala	Asn	Ala	Ala 365	Gly	Glu	Ala
Thr Ala 370	Thr Val	Glu Val	Ser 375	Ile	Val	Gln	Leu	Pro 380	His	Leu	Ser	Asn
Ser Thr 385	Ser Arg	Thr Ala		Pro	Lys	Ser	Arg 395	Leu	Ser	Asp	Ile	Thr 400
Gly Ser	Ser Lys	Thr Ser	Arg	Gly	Gly	Gly 410	Gly	Ser	Gly	Gly	Gly 415	Glu
Pro Pro	Lys Ser 420	Pro Pro	Glu	Arg	Ala 425	Val	Leu	Val	Ser	Glu 430	Val	Thr
Thr Thr	Ser Ala 435	Leu Ala	Lys	Trp 440	Ser	Val	Ser	Lys	Ser 445	Thr	Pro	Arg
Val Lys 450	Met Tyr	Gln Leu	Gln 455	Tyr	Asn	Cys	Ser	Asp 460	Asp	Glu	Val	Leu
Ile Tyr 465	Arg Met	Ile Pro		Ser	Asn	Lys	Ala 475	Phe	Val	Val	Asn	Asn 480
Leu Val	Ser Gly	Thr Gly 485	Tyr	Asp	Leu	Cys 490	Val	Leu	Ala	Met	Trp 495	Asp
Asp Thr	Ala Thr 500	Thr Leu	Thr	Ala	Thr 505	Asn	Ile	Val	Gly	Cys 510	Ala	Gln
Phe Phe	Thr Lys 515	Ala Asp	Tyr	Pro 520	Gln	Cys	Gln	Ser	Met 525	His	Ser	Gln
Ile Leu 530	Gly Gly	Thr Met	Ile 535	Leu	Val	Ile	Gly	Gly 540	Ile	Ile	Val	Ala
Thr Leu 545	Leu Val	Phe Ile		Ile	Leu	Met	Val 555	Arg	Tyr	Lys	Val	Cys 560

Asn His Glu Ala Pro Ser Lys Met Ala Ala Ala Val Ser Asn Val Tyr 565 570 575

Ser Gln Thr Asn Gly Ala Gln Pro Pro Pro Pro Ser Ser Ala Pro Ala 580 585 590

Gly Ala Pro Pro Gln Gly Pro Pro Lys Val Val Arg Asn Glu Leu 595 600 605

Leu Asp Phe Thr Ala Ser Leu Ala Arg Ala Ser Asp Ser Ser Ser Ser 610 620

Ser Ser Leu Gly Ser Gly Glu Ala Ala Gly Leu Gly Arg Ala Pro Trp 625 630 635

Arg Leu Pro Pro Ser Ala Pro Arg Pro Lys Pro Ser Leu Asp Arg Leu 645 650 655

Met Gly Ala Phe Ala Ser Leu Asp Leu Lys Ser Gln Arg Lys Glu Glu 660 665 670

Leu Leu Asp Ser Arg Thr Pro Ala Gly Arg Gly Ala Gly Thr Ser Ala 675 680 685

Arg Gly His His Ser Asp Arg Glu Pro Leu Leu Gly Pro Pro Ala Ala 690 695 700

Arg Ala Arg Ser Leu Leu Pro Leu Pro Leu Glu Gly Lys Ala Lys Arg 705 710 715 720

Ser His Ser Phe Asp Met Gly Asp Phe Ala Ala Ala Ala Ala Gly Gly 725 730 735

Val Val Pro Gly Gly Tyr Ser Pro Pro Arg Arg Val Ser Asn Ile Trp
740 745 750

Thr Lys Arg Ser Leu Ser Val Asn Gly Met Leu Leu Pro Phe Glu Glu
755 760 765

Ser Asp Leu Val Gly Ala Arg Gly Thr Phe Gly Ser Ser Glu Trp Val 770 780

Met Glu Ser Thr Val 785

<210> 43

<211> 788

<212> PRT

<213> Mus musculus

<400> 43

Met Glu Thr Leu Leu Gly Gly Leu Leu Ala Phe Gly Met Ala Phe Ala 1 5 10 15

Val Val Asp Ala Cys Pro Lys Tyr Cys Val Cys Gln Asn Leu Ser Glu 20 25 30

Ser Leu Gly Thr Leu Cys Pro Ser Lys Arg Leu Leu Phe Val Pro Pro 35 40 45

Asp Ile Asp Arg Arg Thr Val Glu Leu Arg Leu Gly Gly Asn Phe Ile 50 55 60

Ile His Ile Gly Arg Gln Asp Phe Ala Asn Met Thr Gly Leu Val Asp 65 70 75 80

Leu Thr Leu Ser Arg Asn Thr Ile Ser His Ile Gln Pro Phe Ser Phe 85 90 95

Leu Asp Leu Glu Ser Leu Arg Ser Leu His Leu Asp Ser Asn Arg Leu
100 105 110

Pro Ser Leu Gly Glu Asp Thr Leu Arg Gly Leu Val Asn Leu Gln His 115 120 125

Leu Ile Val Asn Asn Asn Gln Leu Gly Gly Ile Ala Asp Asp Ala Phe 130 135 140

Leu His Gly Leu Pro Trp Asp Ser Val Arg Arg Met Val Asn Leu His
165 170 175

Gln Leu Ser Leu Asp His Asn Leu Leu Asp His Ile Ala Glu Gly Thr 180 185 190

Phe Ala Asp Leu Gln Lys Leu Ala Arg Leu Asp Leu Thr Ser Asn Arg 195 200 205

Leu Gln Lys Leu Pro Pro Asp Pro Ile Phe Ala Arg Ser Gln Ala Ser 210 215 220

Leu Leu Thr Ala Thr Pro Phe Ala Pro Pro Leu Ser Phe Ser Phe Gly

Gly Asn Pro Leu His Cys Asn Cys Glu Leu Leu Trp Leu Arg Arg Leu 245 250 255

Glu Arg Asp Asp Leu Glu Thr Cys Gly Ser Pro Gly Ser Leu Lys 260 265 270

Gly Arg Tyr Phe Trp His Ile Arg Glu Glu Glu Phe Val Cys Glu Pro 275 280 285

Pro Leu Ile Thr Gln His Thr His Lys Leu Leu Val Leu Glu Gly Gln 290 295 300

Ala Ala Thr Leu Lys Cys Lys Ala Ile Gly Asp Pro Ser Pro Leu Ile 305 310 315 320

His Trp Val Ala Pro Asp Asp Arg Leu Val Gly Asn Ser Ser Arg Thr 325 330 335

Ala Val Tyr Asp Asn Gly Thr Leu Asp Ile Leu Ile Thr Thr Ser Gln 340 345 350

Asp Ser Gly Pro Phe Thr Cys Ile Ala Ala Asn Ala Ala Gly Glu Ala 355 360 365

Thr Ala Thr Val Glu Val Ser Ile Val Gln Leu Pro His Leu Ser Asn 370 380

Ser Thr Ser Arg Met Ala Pro Pro Lys Ser Arg Leu Ser Asp Ile Thr 385 390 395 400

Gly Ser Ser Lys Thr Ser Arg Gly Gly Gly Gly Ser Gly Ala Gly Glu 405 410 415

Pro Pro Lys Ser Thr Pro Glu Arg Ala Val Leu Val Ser Asp Val Thr 420 425 430

Thr Thr Ser Ala Leu Val Lys Trp Ser Val Ser Lys Ser Ala Pro Arg
435 440 445

Val Lys Met Tyr Gln Leu Gln Tyr Asn Cys Ser Asp Asp Glu Val Leu 450 455 460

Ile Tyr Arg Met Ile Pro Ala Ser Asn Lys Ala Phe Val Val Asn Asn 465 470 475 480

Leu Val Ser Gly Thr Gly Tyr Asp Leu Cys Val Leu Ala Met Trp Asp

Asp	Thr	Ala	Thr 500	Thr	Leu	Thr	Ala	Thr 505	Asn	Ile	Val	Gly	Cys 510	Ala	Gln
Phe	Phe	Thr 515	Lys	Ala	Asp	Tyr	Pro 520	Gln	Cys	Gln	Ser	Met 525	His	Ser	Gln
Ile	Lys 530	Gly	Gly	Thr	Met	Ile 535	Leu	Val	Ile	Gly	Gly 540	Ile	Ile	Val	Ala
Thr 545	Leu	Leu	Val	Phe	Ile 550	Val	Ile	Leu	Met	Val 555	Arg	Tyr	Lys	Val	Cys 560
Asn	His	Asp	Thr	Pro 565	Gly	Lys	Met	Ala	Ala 570	Ala	Thr	Val	Ser	Asn 575	Val
Tyr	Ser	Gln	Thr 580	Asn	Gly	Ser	Gln	Pro 585	Pro	Pro	Leu	Gly	Gly 590	Ile	Pro
Val	Gly	Gln 595	Leu	Pro	Gln	Ala	Pro 600	Pro	Lys	Val	Val	Val 605	Arg	Asn	Glu
Leu	Met 610	Asp	Phe	Ser	Thr	Ser 615	Leu	Ala	Arg	Ala	Cys 620	Asp	Ser	Ser	Ser
Ser 625	Ser	Ser	Leu	Gly	Ser 630	Gly	Glu	Ala	Ala	Gly 635	Leu	Gly	Arg	Gly	Pro 640
Trp	Arg	Leu	Pro	Pro 645	Pro	Ala	Pro	Arg	Pro 650	Lys	Pro	Ser	Leu	Asp 655	Arg
Leu	Met	Gly	Ala 660	Phe	Ala	Ser	Leu	Asp 665	Leu	Lys	Ser	Gln	Arg 670	Lys	Glu
Glu	Leu	Leu 675	Asp	Ser	Arg	Thr	Pro 680	Ala	Gly	Arg	Gly	Ala 685	Gly	Thr	Ser
Ser	Arg 690	Gly	His	His	Ser	Asp 695	Arg	Glu	Pro	Leu	Leu 700	Gly	Pro	Pro	Ala
Thr 705	Arg	Ala	Arg	Ser	Leu 710	Leu	Pro	Leu	Pro	Leu 715	Glu	Gly	Lys	Ala	Lys 720
Arg	Ser	His	Ser	Phe 725	Asp	Met	Gly	Asp	Phe 730	Ala	Ala	Ala	Ala	Ala 735	Ala

Val Pro Gly Gly Tyr Ser Pro Pro Arg Arg Val Ser Asn Ile Trp Thr

740 745 750

Lys Arg Ser Leu Ser Val Asn Gly Met Leu Leu Pro Phe Glu Glu Ser 755 760 765

Asp Leu Val Gly Ala Arg Gly Thr Phe Gly Ser Ser Glu Trp Val Met 770 780

Glu Ser Thr Val 785

<210> 44

<211> 25

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: LRR, Leucine Rich Repeat domain sequence

<400> 44

Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Ser Gly Ser Leu 1 5 10 15

Pro Pro Glu Ser Phe Gly Asn Leu Pro 20 25

<210> 45

<211> 24

<212> PRT

<213> Homo sapiens

<400> 45

Arg Val Val Glu Leu Arg Leu Thr Asp Asn Phe Ile Ala Ala Val Arg

1 5 10 15

<210> 46

<211> 25

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: LRR, Leucine Rich Repeat domain sequence <400> 46 Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Ser Gly Ser Leu 5 10 Pro Pro Glu Ser Phe Gly Asn Leu Pro 20 <210> 47 <211> 24 <212> PRT <213> Homo sapiens <400> 47 Ser Leu Val His Leu Thr Leu Ser Arg Asn Thr Ile Gly Gln Val Ala 5 10 Ala Gly Ala Phe Ala Asp Leu Arg 20 <210> 48 <211> 25 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: LRR, Leucine Rich Repeat domain sequence <400> 48 Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Ser Gly Ser Leu 1. 5 10 15 Pro Pro Glu Ser Phe Gly Asn Leu Pro 20 25 <210> 49 <211> 24 <212> PRT <213> Homo sapiens <400> 49 Ala Leu Arg Ala Leu His Leu Asp Ser Asn Arg Leu Ala Glu Val Arg 1

5

10

15

Gly Asp Gln Leu Arg Gly Leu Gly 20

<210> 50

<211> 25

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: LRR, Leucine Rich Repeat domain sequence

<400> 50

Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Ser Gly Ser Leu

1 5 10 15

Pro Pro Glu Ser Phe Gly Asn Leu Pro 20 25

<210> 51

<211> 24

<212> PRT

<213> Homo sapiens

<400> 51

Asn Leu Arg His Leu Ile Leu Gly Asn Asn Gln Ile Arg Arg Val Glu
1 5 10 15

Ser Ala Ala Phe Asp Ala Phe Leu 20

<210> 52

<211> 25

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: LRR, Leucine Rich Repeat domain sequence

<400> 52

Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Ser Gly Ser Leu

1 5 10 15

20 <210> 53 <211> 24 <212> PRT <213> Homo sapiens <400> 53 Thr Val Glu Asp Leu Asp Leu Ser Tyr Asn Asn Leu Glu Ala Leu Pro 5 Trp Glu Ala Val Gly Gln Met Val 20 <210> 54 <211> 25 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: LRR, Leucine Rich Repeat domain sequence <400> 54 Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Ser Gly Ser Leu 5 10 Pro Pro Glu Ser Phe Gly Asn Leu Pro 20 25 <210> 55 <211> 24 <212> PRT <213> Homo sapiens <400> 55 Asn Leu Asn Thr Leu Thr Leu Asp His Asn Leu Ile Asp His Ile Ala 1 5 10 15

Pro Pro Glu Ser Phe Gly Asn Leu Pro

Glu Gly Thr Phe Val Gln Leu His 20

<211> 25 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: LRR, Leucine Rich Repeat domain sequence <400> 56 Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Ser Gly Ser Leu 10 Pro Pro Glu Ser Phe Gly Asn Leu Pro 20 <210> 57 <211> 23 <212> PRT <213> Homo sapiens <400> 57 Lys Leu Val Arg Leu Asp Met Thr Ser Asn Arg Leu His Lys Leu Pro . 10 Pro Asp Gly Leu Phe Leu Arg 20 <210> 58 <211> 54 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: LRR, Leucine Rich Repeat domain sequence <400> 58 Asn Pro Phe Asn Cys Asp Cys Glu Leu Arg Trp Leu Leu Arg Trp Leu Arg Glu Thr Asn Pro Arg Arg Leu Glu Asp Gln Glu Asp Leu Arg Cys 25 Ala Ser Pro Glu Ser Leu Arg Gly Gln Pro Leu Leu Glu Leu Leu Pro

<210> 56

40

35

50 <210> 59 <211> 46 <212> PRT <213> Homo sapiens <400> 59 Asn Pro Leu His Cys Asn Cys Glu Leu Leu Trp Leu Arg Arg Leu Thr 1 5 10 Arg Glu Asp Asp Leu Glu Thr Cys Ala Thr Pro Glu His Leu Thr Asp 20 25 Arg Tyr Phe Trp Ser Ile Pro Glu Glu Glu Phe Leu Cys Glu 35 40 <210> 60 <211> 45 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Immunoglobin domain sequence Gly Glu Ser Val Thr Leu Thr Cys Ser Val Ser Gly Phe Gly Pro Pro 1 5 10 15 Pro Val Thr Trp Leu Arg Asn Gly Lys Leu Ser Leu Thr Ile Ser Val 20 Thr Pro Glu Asp Ser Gly Gly Thr Tyr Thr Cys Val Val 35 40 45 <210> 61 <211> 59 <212> PRT

Ser Asp Phe Ser Cys Pro

<213> Homo sapiens

<400> 61

Gly Gln Ala Val Ser Leu Arg Cys Arg Ala Val Gly Asp Pro Glu Pro

1 5 10 15

Val Val His Trp Val Ala Pro Asp Gly Arg Leu Leu Gly Asn Ser Ser 20 25 30

Arg Thr Arg Val Arg Gly Asp Gly Thr Leu Asp Val Thr Ile Thr Thr 35 40 45

Leu Arg Asp Ser Gly Thr Phe Thr Cys Ile Ala 50 55

<210> 62

<211> 84

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Fibronectin Type III domain sequence

<400> 62

Pro Ser Ala Pro Thr Asn Leu Thr Val Thr Asp Val Thr Ser Thr Ser 1 5 10 15

Leu Thr Leu Ser Trp Ser Pro Pro Thr Gly Asn Gly Pro Ile Thr Gly
20 25 30

Tyr Glu Val Thr Tyr Arg Gln Pro Lys Asn Gly Gly Glu Trp Asn Glu 35 40 45

Leu Thr Val Pro Gly Thr Thr Thr Ser Tyr Thr Leu Thr Gly Leu Lys 50 55 60

Pro Gly Thr Glu Tyr Glu Val Arg Val Gln Ala Val Asn Gly Gly Gly 65 70 75 80

Gly Pro Glu Ser

<210> 63

<211> 81

<212> PRT

<213> Homo sapiens

<400> 63

Ser Ala Ala Glu Arg Arg Leu Val Ala Ala Glu Leu Thr Ser Asn Ser

1 5 10 15

Val Leu Ile Arg Trp Pro Ala Gln Arg Pro Val Pro Gly Ile Arg Met
20 25 30

Tyr Gln Val Gln Tyr Asn Ser Ser Val Asp Asp Ser Leu Val Tyr Arg
35 40 45

Met Ile Pro Ser Thr Ser Gln Thr Phe Leu Val Asn Asp Leu Ala Ala 50 55 60

Gly Arg Ala Tyr Asp Leu Cys Val Leu Ala Val Tyr Asp Asp Gly Ala 65 70 75 80

Thr

<210> 64

<211> 405

<212> PRT

<213> Homo sapiens

<400> 64

Met Glu Arg Leu Gln Lys Gln Pro Leu Thr Ser Pro Gly Ser Val Ser 1 5 10 15

Pro Ser Arg Asp Ser Ser Val Pro Gly Ser Pro Ser Ser Ile Val Ala 20 25 30

Lys Met Asp Asn Gln Val Leu Gly Tyr Lys Asp Leu Ala Ala Ile Pro 35 40 45

Lys Asp Lys Ala Ile Leu Asp Ile Glu Arg Pro Asp Leu Met Ile Tyr 50 55 60

Glu Pro His Phe Thr Tyr Ser Leu Leu Glu His Val Glu Leu Pro Arg
65 70 75 80

Ser Arg Glu Arg Ser Leu Ser Pro Lys Ser Thr Ser Pro Pro Pro Ser 85 90 95

Pro Glu Val Trp Ala Asp Ser Arg Ser Pro Gly Ile Ile Ser Gln Ala 100 105 110

Ser Ala Pro Arg Thr Thr Gly Thr Pro Arg Thr Ser Leu Pro His Phe 115 120 125

His	His 130	Pro	Glu	Thr	Ser	Arg 135	Pro	Asp	Ser	Asn	Ile 140	Tyr	Lys	Lys	Pro
Pro 145	Ile	Tyr	Lys	Gln	Arg 150	Glu	Ser	Val	Gly	Gly 155	Ser	Pro	Gln	Thr	Lys 160
His	Leu	Ile	Glu	Asp 165	Leu	Ile	Ile	Glu	Ser 170	Ser	Lys	Phe	Pro	Ala 175	Ala
Gln	Pro	Pro	Asp 180	Pro	Asn	Gln	Pro	Ala 185	Lys	Ile	Glu	Thr	Asp 190	Tyr	Trp
Pro	Cys	Pro 195	Pro	Ser	Leu	Ala	Val 200	Val	Glu	Thr	Glu	Trp 205	Arg	Lys	Arg
Lys	Ala 210	Ser	Arg	Arg	Gly	Ala 215	Glu	Glu	Glu	Glu	Glu 220	Glu	Glu	Asp	Asp
Asp 225	Ser	Gly	Glu	Glu	Met 230	Lys	Ala	Leu	Arg	Glu 235	Arg	Gln	Arg	Glu	Glu 240
Leu	Ser	Lys	Val	Thr 245	Ser	Asn	Leu	Gly	Lys 250	Met	Ile	Leu	Lys	Glu 255	Glu
Met	Glu	Lys	Ser 260	Leu	Pro	Ile	Arg	Arg 265	Lys	Thr	Arg	Ser	Leu 270	Pro	Asp
Arg	Thr	Pro 275	Phe	His	Thr	Ser	Leu 280	His	Gln	Gly	Thr	Ser 285	Lys	Ser	Ser
Ser	Leu 290	Pro	Ala	Tyr	Gly	Arg 295	Thr	Thr	Leu	Ser	Arg 300	Leu	Gln	Ser	Thr
Glu 305	Phe	Ser	Pro	Ser	Gly 310	Ser	Glu	Thr	Gly	Ser 315	Pro	Gly	Leu	Gln	Asn 320
Gly	Glu	Gly	Gln	Arg 325	Gly	Arg	Met	Asp	Arg 330	Gly	Asn	Ser	Leu	Pro 335	Cys
Val	Leu	Glu	Gln 340	Lys	Ile	Tyr	Pro	Tyr 345	Glu	Met	Leu	Val	Val 350	Thr	Asn
Lys	Gly	Arg 355	Thr	Lys	Leu	Pro	Pro 360	Gly	Val	Asp	Arg	Met 365	Arg	Leu	Glu
Arg	His 370	Leu	Ser	Ala	Glu	Asp 375	Phe	Ser	Arg	Val	Phe 380	Ala	Met	Ser	Pro

Glu Glu Phe Gly Lys Leu Ala Leu Trp Lys Arg Asn Glu Leu Lys Lys 385 390 395 400

Lys Ala Ser Leu Phe 405

<210> 65

<211> 383

<212> PRT

<213> Homo sapiens

<400> 65

Met Glu Arg Leu Gln Lys Gln Pro Leu Thr Ser Pro Gly Ser Val Ser 1 5 10 15

Pro Ser Arg Asp Ser Ser Val Pro Gly Ser Pro Ser Ser Ile Val Ala 20 25 30

Lys Met Asp Asn Gln Val Leu Gly Tyr Lys Asp Leu Ala Ala Ile Pro 35 40 45

Lys Asp Lys Ala Ile Leu Asp Ile Glu Arg Pro Asp Leu Met Ile Tyr 50 55 60

Glu Pro His Phe Thr Tyr Ser Leu Leu Glu His Val Glu Leu Pro Arg
65 70 75 80

Ser Arg Glu Arg Ser Leu Ser Pro Lys Ser Thr Ser Pro Pro Pro Ser 85 90 95

Pro Glu Val Trp Ala Asp Ser Arg Ser Pro Gly Ile Ile Ser Gln Ala 100 105 110

Ser Ala Pro Arg Thr Thr Gly Thr Pro Arg Thr Ser Leu Pro His Phe 115 120 125

His His Pro Glu Thr Ser Arg Pro Asp Ser Asn Ile Tyr Lys Lys Pro 130 135 140

Pro Ile Tyr Lys Gln Arg Glu Ser Val Gly Gly Ser Pro Gln Thr Lys 145 150 155 160

His Leu Ile Glu Asp Leu Ile Ile Glu Ser Ser Lys Phe Pro Ala Ala 165 170 175

Gln Pro Pro Asp Pro Asn Gln Pro Ala Lys Ile Glu Thr Asp Tyr Trp 180 185 190

Asp Ser Gly Glu Glu Met Lys Ala Leu Arg Glu Arg Gln-Arg Glu Glu 225 230 235 240

Leu Ser Lys Val Thr Ser Asn Leu Gly Lys Met Ile Leu Lys Glu Glu 245 250 255

Met Glu Lys Ser Leu Pro Ile Arg Arg Lys Thr Arg Ser Leu Pro Asp $260 \hspace{1.5cm} 265 \hspace{1.5cm} 270 \hspace{1.5cm}$

Arg Thr Pro Phe His Thr Ser Leu His Gln Gly Thr Ser Lys Ser Ser 275 280 285

Ser Leu Pro Ala Tyr Gly Arg Thr Thr Leu Ser Arg Leu Gln Ser Thr 290 295 300

Glu Phe Ser Pro Ser Gly Ser Glu Thr Gly Ser Pro Asp Pro Gln Ile 305 310 315

Tyr Pro Tyr Glu Met Leu Val Val Thr Asn Lys Gly Arg Thr Lys Leu 325 330 335

Pro Pro Gly Val Asp Arg Met Arg Leu Glu Arg His Leu Ser Ala Glu 340 345 350

Asp Phe Ser Arg Val Phe Ala Met Ser Pro Glu Glu Phe Gly Lys Leu 355 360 365

Ala Leu Trp Lys Arg Asn Glu Leu Lys Lys Lys Ala Ser Leu Phe 370 375 380

<210> 66

<211> 383

<212> PRT

<213> Homo sapiens

<400> 66

Met Glu Arg Leu Gln Lys Gln Pro Leu Thr Ser Pro Gly Ser Val Ser 1 5 10 15

Pro Ser Arg Asp Ser Ser Val Pro Gly Ser Pro Ser Ser Ile Val Ala

Lys	Met	Asp 35	Asn	Gln	Val	Leu	Gly 40	Tyr	Lys	Asp	Leu	Ala 45	Ala	Ile	Pro
Lys	Asp 50	Lys	Ala	Ile	Leu	Asp 55	Ile	Glu	Arg	Pro	Asp 60	Leu	Met	Ile	Tyr
Glu 65	Pro	His	Phe	Thr	Tyr 70	Ser	Leu	Leu	Glu	His 75	Val	Glu	Leu	Pro	Arg 80
Gln	Arg	Glu	Arg	Ser 85	Leu	Ser	Pro	Lys	Ser 90	Thr	Ser	Pro	Pro	Pro 95	Ser
Pro	Glu	Val	Trp 100	Ala	Asp	Ser	Arg	Ser 105	Pro	Gly	Ile	Ile	Ser 110	Gln	Ala
Ser	Ala	Pro 115	Arg	Thr	Thr	Gly	Thr 120	Pro	Arg	Thr	Ser	Leu 125	Pro	His	Phe
His	His 130	Pro	Glu	Thr	Ser	Arg 135	Pro	Asp	Ser	Asn	Ile 140	Tyr	Lys	Lys	Pro
Pro 145	Ile	Tyr	Lys	Gln	Arg 150	Glu	Ser	Val	Gly	Gly 155	Ser	Pro	Gln	Thr	Lys 160
His	Leu	Ile	Glu	Asp 165	Leu	Ile	Ile	Glu	Ser 170	Ser	Lys	Phe	Pro	Ala 175	Ala
Gln	Pro	Pro	Asp 180	Pro	Asn	Gln	Pro	Ala 185	Lys	Ile	Glu	Thr	Asp 190	Tyr	Trp
Pro	Cys	Pro 195	Pro	Ser	Leu	Ala	Val 200	Val	Glu	Thr	Glu	Trp 205	Arg	Lys	Arg
Lys	Ala 210	Ser	Arg	Arg	Gly	Ala 215	Glu	Glu	Glu	Glu	Glu 220	Glu	Glu	Asp	Asp
Asp 225	Ser	Gly	Glu	Glu	Met 230	Lys	Ala	Leu	Arg	Glu 235	Arg	Gln	Arg	Glu	Glu 240
Leu	Ser	Lys	Val	Thr 245	Ser	Asn	Leu	Gly	Lys 250	Met	Ile	Leu	Lys	Glu 255	Glu
Met	Glu	Lys	Ser 260	Leu	Pro	Ile	Arg	Arg 265	Lys	Thr	Arg	Ser	Leu 270	Pro	Asp

Arg Thr Pro Phe His Thr Ser Leu His Gln Gly Thr Ser Lys Ser Ser

Ser Leu Pro Arg Tyr Gly Arg Thr Thr Leu Ser Arg Leu Gln Ser Thr 290 295 300

Glu Phe Ser Pro Ser Gly Ser Glu Thr Gly Ser Pro Gly Leu Gln Ile 305 310 315 320

Tyr Pro Tyr Glu Val Leu Val Val Thr Asn Lys Gly Arg Thr Lys Leu 325 330 335

Pro Pro Gly Val Asp Arg Met Arg Leu Glu Arg His Leu Ser Ala Glu
340 345 350

Asp Phe Ser Arg Val Ser Ala Met Ser Pro Glu Glu Phe Gly Lys Leu 355 360 365

Ala Leu Trp Lys Arg Asn Glu Leu Lys Lys Lys Ala Ser Leu Phe 370 375 380

<210> 67

<211> 405

<212> PRT

<213> Homo sapiens

<400> 67

Met Glu Arg Leu Gln Lys Gln Pro Leu Thr Ser Pro Gly Ser Val Ser

1 5 10 15

Pro Ser Arg Asp Ser Ser Val Pro Gly Ser Pro Ser Ser Ile Val Ala 20 25 30

Lys Met Asp Asn Gln Val Leu Gly Tyr Lys Asp Leu Ala Ala Ile Pro 35 40 45

Lys Asp Lys Ala Ile Leu Asp Ile Glu Arg Pro Asp Leu Met Ile Tyr 50 55 60

Glu Pro His Phe Thr Tyr Ser Leu Leu Glu His Val Glu Leu Pro Arg
65 70 75 80

Gln Arg Glu Arg Ser Leu Ser Pro Lys Ser Thr Ser Pro Pro Pro Ser 85 90 95

Pro Glu Val Trp Ala Asp Ser Arg Ser Pro Gly Ile Ile Ser Gln Ala 100 105 110

Ser	Ala	Pro 115	Arg	Thr	Thr	Gly	Thr 120	Pro	Arg	Thr	Ser	Leu 125	Pro	His	Phe
His	His 130	Pro	Glu	Thr	Ser	Arg 135	Pro	Asp	Ser	Asn	Ile 140	Tyr	Lys	Lys	Pro
Pro 145	Ile	Tyr	Lys	Gln	Arg 150	Glu	Ser	Val	Gly	Gly 155	Ser	Pro	Gln	Thr	Lys 160
His	Leu	Ile	Glu	Asp 165	Leu	Ile	Ile	Glu	Ser 170	Ser	Lys	Phe	Pro	Ala 175	Ala
Gln	Pro	Pro	Asp 180	Pro	Asn	Gln	Pro	Ala 185	Lys	Ile	Glu	Thr	Asp 190	Tyr	Trp
Pro	Cys	Pro 195	Pro	Ser	Leu	Ala	Val 200	Val	Glu	Thr	Glu	Trp 205	Arg	Lys	Arg
Lys	Ala 210	Ser	Arg	Arg	Gly	Ala 215	Glu	Glu	Glu	Glu	Glu 220	Glu	Glu	Asp	Asp
Asp 225	Ser	Gly	Glu	Glu	Met 230	Lys	Ala	Leu	Arg	Glu 235	Arg	Gln	Arg	Glu	Glu 240
Leu	Ser	Lys	Val	Thr 245	Ser	Asn	Leu	Gly	Lys 250	Met	Ile	Leu	Lys	Glu 255	Glu
Met	Glu	Lys	Ser 260	Leu	Pro	Ile	Arg	Arg 265	Lys	Thr	Arg	Ser	Leu 270	Pro	Asp
Arg	Thr	Pro 275	Phe	His	Thr	Ser	Leu 280	His	Gln	Gly	Thr	Ser 285	Lys	Ser	Ser
Ser	Leu 290	Pro	Arg	Tyr	Gly	Arg 295	Thr	Thr	Leu	Ser	Arg 300	Leu	Gln	Ser	Thr
Glu 305	Phe	Ser	Pro	Ser	Gly 310	Ser	Glu	Thr	Gly	Ser 315	Pro	Gly	Leu	Gln	Asn 320
Gly	Glu	Gly	Gln	Arg 325	Gly	Arg	Met	Asp	Arg 330	Gly	Asn	Ser	Leu	Pro 335	Cys
Val	Leu	Glu	Gln 340	Lys	Ile	Tyr	Pro	Tyr 345	Glu	Met	Leu	Val	Val 350	Thr	Asn
Lys	Gly	Arg 355	Thr	Lys	Leu	Pro	Pro 360	Gly	Val	Asp	Arg	Met 365	Arg	Leu	Glu

Arg His Leu Ser Ala Glu Asp Phe Ser Arg Val Phe Ala Met Ser Pro 370 380

Glu Glu Phe Gly Lys Leu Ala Leu Trp Lys Arg Asn Glu Leu Lys Lys 385 390 395 400

Lys Ala Ser Leu Phe 405

<210> 68

<211> 405

<212> PRT

<213> Homo sapiens

<400> 68

Met Glu Arg Leu Gln Lys Gln Pro Leu Thr Ser Pro Gly Ser Val Ser 1 5 10 15

Pro Ser Arg Asp Ser Ser Val Pro Gly Ser Pro Ser Ser Ile Val Ala
20 25 30

Lys Met Asp Asn Gln Val Leu Gly Tyr Lys Asp Leu Ala Ala Ile Pro 35 40 45

Lys Asp Lys Ala Ile Leu Asp Ile Glu Arg Pro Asp Leu Met Ile Tyr 50 55 60

Glu Pro His Phe Thr Tyr Ser Leu Leu Glu His Val Glu Leu Pro Arg
65 70 75 80

Ser Arg Glu Arg Ser Leu Ser Pro Lys Ser Thr Ser Pro Pro Pro Ser 85 90 95

Pro Glu Val Trp Ala Asp Ser Arg Ser Pro Gly Ile Ile Ser Gln Ala 100 105 110

Ser Ala Pro Arg Thr Thr Gly Thr Pro Arg Thr Ser Leu Pro His Phe 115 120 125

His His Pro Glu Thr Ser Arg Pro Asp Ser Asn Ile Tyr Lys Lys Pro 130 135 140

Pro Ile Tyr Lys Gln Arg Glu Ser Val Gly Gly Ser Pro Gln Thr Lys 145 150 155 160

His Leu Ile Glu Asp Leu Ile Ile Glu Ser Ser Lys Phe Pro Ala Ala 165 170 175

Gln Pro Pro Asp Pro Asn Gln Pro Ala Lys Ile Glu Thr Asp Tyr Trp Pro Cys Pro Pro Ser Leu Ala Val Val Glu Thr Glu Trp Arg Lys Arg Lys Ala Ser Arg Arg Gly Ala Glu Glu Glu Glu Glu Glu Asp Asp Asp Ser Gly Glu Glu Met Lys Ala Leu Arg Glu Arg Gln Arg Glu Glu Leu Ser Lys Val Thr Ser Asn Leu Gly Lys Met Ile Leu Lys Glu Glu Met Glu Lys Ser Leu Pro Ile Arg Arg Lys Thr Arg Ser Leu Pro Asp Arg Thr Pro Phe His Thr Ser Leu His Gln Gly Thr Ser Lys Ser Ser Ser Leu Pro Ala Tyr Gly Arg Thr Thr Leu Ser Arg Leu Gln Ser Thr Glu Phe Ser Pro Ser Gly Ser Glu Thr Gly Ser Pro Ala Pro Gln Asn Gly Glu Gly Gln Arg Gly Arg Met Asp Arg Gly Asn Ser Leu Pro Cys Val Leu Glu Gln Lys Ile Tyr Pro Tyr Glu Met Leu Val Val Thr Asn Lys Gly Arg Thr Lys Leu Pro Pro Gly Val Asp Arg Met Arg Leu Glu Arg His Leu Ser Ala Glu Asp Phe Ser Arg Val Phe Ala Met Ser Pro Glu Glu Phe Gly Lys Leu Ala Leu Trp Lys Arg Asn Glu Leu Lys Lys

<210> 69

Lys Ala Ser Leu Phe

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<220>
<223> Description of Artificial Sequence: VHP, Villin
      headpiece domain sequence
<400> 69
Tyr Leu Ser Asp Glu Asp Phe Glu Glu Val Phe Gly Met Thr Lys Glu
                                     10
Glu Phe Tyr Lys Leu Pro Leu Trp Lys Gln Asn Gln Leu Lys Lys
                                                      30
                                 25
             20
Leu Gly Leu Phe
         35
<210> 70
<211> 36
<212> PRT
<213> Homo sapiens
<400> 70
His Leu Ser Ala Glu Asp Phe Ser Arg Val Phe Ala Met Ser Pro Glu
Glu Phe Gly Lys Leu Ala Leu Trp Lys Arg Asn Glu Leu Lys Lys
                                 25
Ala Ser Leu Phe
         35
<210> 71
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: VHP, Villin
      headpiece domain sequence
<400> 71
Tyr Leu Ser Asp Glu Asp Phe Glu Glu Val Phe Gly Met Thr Lys Glu
  1
                  5
                                     10
                                                          15
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<211> 36 <212> PRT

<213> Artificial Sequence

Glu Phe Tyr Lys Leu Pro Ala Trp Lys Gln Asn Gln Leu Lys Lys 20 25 30

Leu Gly Leu Phe 35

<210> 72

<211> 36

<212> PRT

<213> Homo sapiens

<400> 72

His Leu Ser Ala Glu Asp Phe Ser Arg Val Phe Ala Met Ser Pro Glu
1 5 10 15

Glu Phe Gly Lys Leu Ala Leu Trp Lys Arg Asn Glu Leu Lys Lys 20 25 30

Ala Ser Leu Phe 35

<210> 73

<211> 959

<212> PRT

<213> Homo sapiens

<400> 73

Met Glu Lys Met Leu Ala Gly Cys Phe Leu Leu Ile Leu Gly Gln Ile 1 5 10 15

Val Leu Leu Pro Cys Glu Ala Arg Glu Arg Ser Arg Gly Arg Ser Ile 20 25 30

Ser Arg Gly Arg His Ala Arg Thr His Pro Gln Thr Ala Leu Leu Glu 35 40 45

Ser Ser Cys Glu Asn Lys Arg Ala Asp Leu Val Phe Ile Ile Asp Ser 50 55 60

Ser Arg Ser Val Asn Thr His Asp Tyr Ala Lys Val Lys Glu Phe Ile 65 70 75 80

Val Asp Ile Leu Gln Phe Leu Asp Ile Gly Pro Asp Val Thr Arg Val
85 90 95

Gly Leu Leu Gln Tyr Gly Ser Thr Val Lys Asn Glu Phe Ser Leu Lys

			100					100					110		
Thr	Phe	Lys 115	Arg	Lys	Ser	Glu	Val 120	Glu	Arg	Ala	Val	Lys 125	Arg	Met	Arg
His	Leu 130	Ser	Thr	Gly	Thr	Met 135	Thr	Gly	Leu	Ala	Ile 140	Gln	Tyr	Ala	Leu
Asn 145	Ile	Ala	Phe	Ser	Glu 150	Ala	Glu	Gly	Ala	Arg 155	Pro	Leu	Arg	Glu	Asn 160
Val	Pro	Arg	Val	Ile 165	Met	Ile	Val	Thr	Asp 170	Gly	Arg	Pro	Gln	Asp 175	Ser
Val	Ala	Glu	Val 180	Ala	Ala	Lys	Ala	Arg 185	Asp	Thr	Gly	Ile	Leu 190	Ile	Phe
Ala	Ile	Gly 195	Val	Gly	Gln	Val	Asp 200	Phe	Asn	Thr	Leu	Lys 205	Ser	Ile	Gly
Ser	Glu 210	Pro	His	Glu	Asp	His 215	Val	Phe	Leu	Val	Ala 220	Asn	Phe	Ser	Gln
Ile 225	Glu	Thr	Leu	Thr	Ser 230	Val	Phe	Gln	Lys	Lys 235	Leu	Cys	Thr	Ala	His 240
Met	Cys	Ser	Thr	Leu 245	Glu	His	Asn	Cys	Ala 250	His	Phe	Cys	Ile	Asn 255	Ile
Pro	Gly	Ser	Tyr 260	Val	Cys	Arg	Cys	Lys 265	Gln	Gly	Tyr	Ile	Leu 270	Asn	Ser
Asp	Gln	Thr 275	Thr	Cys	Arg	Ile	Gln 280	Asp	Leu	Cys	Ala	Met 285	Glu	Asp	His
Asn	Cys 290	Glu	Gln	Leu	Cys	Val 295	Asn	Val	Pro	Gly	Ser 300	Phe	Val	Cys	Glu
Cys 305	Tyr	Ser	Gly	Tyr	Ala 310	Leu	Ala	Glu	Asp	Gly 315	Lys	Arg	Cys	Val	Ala 320
	_	_	_		_		_								

Asn Ala Asp Gly Ser Tyr Leu Cys Gln Cys His Glu Gly Phe Ala Leu 340 345 350

Val Asp Tyr Cys Ala Ser Glu Asn His Gly Cys Glu His Glu Cys Val

Asn Pro Asp Glu Lys Thr Cys Thr Lys Ile Asp Tyr Cys Ala Ser Ser

Asn	His 370	Gly	Суз	Gln	Tyr	Glu 375	Cys	Val	Asn	Thr	Asp 380	Asp	Ser	Tyr	Ser
Cys 385	His	Суѕ	Leu	Lys	Gly 390	Phe	Thr	Leu	Asn	Pro 395	Asp	Lys	Lys	Thr	Cys 400
Arg	Arg	Ile	Asn	Tyr 405	Cys	Ala	Leu	Asn	Lys 410	Pro	Gly	Cys	Glu	His 415	Glu
Cys	Val	Asn	Met 420	Glu	Glu	Ser	Tyr	Tyr 425	Cys	Arg	Cys	His	Arg 430	Gly	Tyr
Thr	Leu	Asp 435	Pro	Asn	Gly	Lys	Pro 440	Cys	Ser	Arg	Val	Asp 445	His	Cys	Ala
Gln	Gln 450	Asp	His	Gly	Cys	Glu 455	Gln	Leu	Cys	Leu	Asn 460	Thr	Glu	Asp	Ser
Phe 465	Val	Cys	Gln	Cys	Ser 470	Glu	Gly	Phe	Leu	Ile 475	Asn	Glu	Asp	Leu	Lys 480
Thr	Cys	Ser	Arg	Val 485	Asp	Tyr	Cys	Leu	Leu 490	Ser	Asp	His	Gly	Cys 495	Glu
Tyr	Ser	Cys	Val 500	Asn	Met	Asp	Arg	Ser 505	Phe	Ala	Cys	Gln	Cys 510	Pro	Glu
Gly	His	Val 515	Leu	Arg	Ser	Asp	Gly 520	Lys	Thr	Cys	Ala	Lys 525	Leu	Asp	Ser
Cys	Ala 530	Leu	Gly	Asp	His	Gly 535	Cys	Glu	His	Ser	Cys 540	Val	Ser	Ser	Glu
Asp 545	Ser	Phe	Val	Cys	Gln 550	Cys	Phe	Glu	Gly	Tyr 555	Ile	Leu	Arg	Glu	Asp 560
Gly	Lys	Thr	Cys	Arg 565	Arg	Lys	Asp	Val	Cys 570	Gln	Ala	Ile	Asp	His 575	Gly
Cys	Glu	His	Ile 580	Cys	Val	Asn	Ser	Asp 585	Asp	Ser	Tyr	Thr	Cys 590	Glu	Cys
Leu	Glu	Gly 595	Phe	Arg	Leu	Thr	Glu 600	Asp	Gly	Lys	Arg	Cys 605	Arg	Ile	Ser

Ser Gly Lys Asp Val Cys Lys Ser Thr His His Gly Cys Glu His Ile

	610					615					620				
Cys 625	Val	Asn	Asn	Gly	Asn 630	Ser	Tyr	Ile	Cys	Lys 635	Cys	Ser	Glu	Gly	Phe 640
Val	Leu	Ala	Glu	Asp 645	Gly	Arg	Arg	Cys	Lys 650	Lys	Cys	Thr	Glu	Gly 655	Pro
Ile	Asp	Leu	Val 660	Phe	Val	Ile	Asp	Gly 665	Ser	Lys	Ser	Leu	Gly 670	Glu	Glu
Asn	Phe	Glu 675	Val	Val	Lys	Gln	Phe 680	Val	Thr	Gly	Ile	Ile 685	Asp	Ser	Leu
Thr	Ile 690	Ser	Pro	Lys	Ala	Ala 695	Arg	Val	Gly	Leu	Leu 700	Gln	Tyr	Ser	Thr
Gln 705	Val	His	Thr	Glu	Phe 710	Thr	Leu	Arg	Asn	Phe 715	Asn	Ser	Ala	Lys	Asp 720
Met	Lys	Lys	Ala	Val 725	Ala	His	Met	Lys	Tyr 730	Met	Gly	Lys	Gly	Ser 735	Met
Thr	Gly	Leu	Ala 740	Leu	Lys	His	Met	Phe 745	Glu	Arg	Ser	Phe	Thr 750	Gln	Gly
Glu	Gly	Ala 755	Arg	Pro	Phe	Ser	Thr 760	Arg	Val	Pro	Arg	Ala 765	Ala	Ile	Val
Phe	Thr 770	Asp	Gly	Arg	Ala	Gln 775	Asp	Asp	Val	Ser	Glu 780	Trp	Ala	Ser	Lys
Ala 785	Lys	Ala	Asn	Gly	Ile 790	Thr	Met	Tyr	Ala	Val 795	Gly	Val	Gly	Lys	Ala 800
Ile	Glu	Glu	Glu	Leu 805	Gln	Glu	Ile	Ala	Ser 810	Glu	Pro	Thr	Asn	Lys 815	His
Leu	Phe	Tyr	Ala 820	Glu	Asp	Phe	Ser	Thr 825	Met	Asp	Glu	Ile	Ser 830	Glu	Lys

Leu Lys Lys Gly Ile Cys Glu Ala Leu Glu Asp Ser Asp Gly Arg Gln

Asp Ser Pro Ala Gly Glu Leu Pro Lys Thr Val Gln Gln Pro Thr Glu

Ser Glu Pro Val Thr Ile Asn Ile Gln Asp Leu Leu Ser Cys Ser Asn

		875	880
865	870		

Phe Ala Val Gln His Arg Tyr Leu Phe Glu Glu Asp Asn Leu Leu Arg 885 890 895

Ser Thr Gln Lys Leu Ser His Ser Thr Lys Pro Ser Gly Ser Pro Leu 900 905 910

Glu Glu Lys His Asp Gln Cys Lys Cys Glu Asn Leu Ile Met Phe Gln 915 920 925

Asn Leu Ala Asn Glu Glu Val Arg Lys Leu Thr Gln Arg Leu Glu Glu 930 935 940

Met Thr Gln Arg Met Glu Ala Leu Glu Asn Arg Leu Arg Tyr Arg 945 950 955

<210> 74

<211> 956

<212> PRT

<213> Homo sapiens

<400> 74

Met Glu Lys Met Leu Ala Gly Cys Phe Leu Leu Ile Leu Gly Gln Ile 1 5 10 15

Val Leu Leu Pro Ala Glu Ala Arg Glu Arg Ser Arg Gly Arg Ser Ile 20 25 30

Ser Arg Gly Arg His Ala Arg Thr His Pro Gln Thr Ala Leu Leu Glu 35 40 45

Ser Ser Cys Glu Asn Lys Arg Ala Asp Leu Val Phe Ile Ile Asp Ser 50 55 60

Ser Arg Ser Val Asn Thr His Asp Tyr Ala Lys Val Lys Glu Phe Ile 65 70 75 80

Val Asp Ile Leu Gln Phe Leu Asp Ile Gly Pro Asp Val Thr Arg Val 85 90 95

Gly Leu Leu Gln Tyr Gly Ser Thr Val Lys Asn Glu Phe Ser Leu Lys 100 105 110

Thr Phe Lys Arg Lys Ser Glu Val Glu Arg Ala Val Lys Arg Met Arg 115 120 125

His Leu Ser Thr Gly Thr Met Thr Gly Leu Ala Ile Gln Tyr Ala Leu Asn Ile Ala Phe Ser Glu Ala Glu Gly Ala Arg Pro Leu Arg Glu Asn Val Pro Arg Val Ile Met Ile Val Thr Asp Gly Arg Pro Gln Asp Ser Val Ala Glu Val Ala Ala Lys Ala Arg Asp Thr Gly Ile Leu Ile Phe Ala Ile Gly Val Gly Gln Val Asp Phe Asn Thr Leu Lys Ser Ile Gly Ser Glu Pro His Glu Asp His Val Phe Leu Val Ala Asn Phe Ser Gln Ile Glu Thr Leu Thr Ser Val Phe Gln Lys Lys Leu Cys Thr Ala His Met Cys Ser Thr Leu Glu His Asn Cys Ala His Phe Cys Ile Asn Ile Pro Gly Ser Tyr Val Cys Arg Cys Lys Gln Gly Tyr Ile Leu Asn Ser Asp Gln Thr Thr Cys Arg Ile Gln Asp Leu Cys Ala Met Glu Asp His Asn Cys Glu Gln Leu Cys Val Asn Val Pro Gly Ser Phe Val Cys Gln Cys Tyr Ser Gly Tyr Ala Leu Ala Glu Asp Gly Lys Arg Cys Val Ala Val Asp Tyr Cys Ala Ser Glu Asn His Gly Cys Glu His Glu Cys Val Asn Ala Asp Gly Ser Tyr Leu Cys Gln Cys His Glu Gly Phe Ala Leu Asn Pro Asp Lys Lys Thr Cys Thr Lys Ile Asp Tyr Cys Ala Ser Ser Asn His Gly Cys Gln His Glu Cys Val Asn Thr Asp Asp Ser Tyr Ser

Cys 385	His	Cys	Leu	Lys	Gly 390	Phe	Thr	Leu	Asn	Pro 395	Asp	Lys	Lys	Thr	Cys 400
Arg	Arg	Ile	Asn	Tyr 405	Cys	Ala	Leu	Asn	Lys 410	Pro	Gly	Cys	Glu	His 415	Glu
Cys	Val	Asn	Met 420	Glu	Glu	Ser	Tyr	Tyr 425	Cys	Arg	Cys	His	Arg 430	Gly	Tyr
Thr	Leu	Asp 435	Pro	Asn	Gly	Lys	Thr 440	Cys	Ser	Arg	Val	Asp 445	His	Cys	Ala
Gln	Gln 450	Asp	His	Gly	Cys	Glu 455	Gln	Leu	Cys	Leu	Asn 460	Thr	Glu	Asp	Ser
Phe 465	Val	Cys	Gln	Cys	Ser 470	Glu	Gly	Phe	Leu	Ile 475	Asn	Glu	Asp	Leu	Lys 480
Thr	Cys	Ser	Arg	Val 485	Asp	Tyr	Cys	Leu	Leu 490	Ser	Asp	His	Gly	Cys 495	Glu
Tyr	Ser	Cys	Val 500	Asn	Met	Asp	Arg	Ser 505	Phe	Ala	Cys	Gln	Cys 510	Pro	Glu
Gly	His	Val 515	Leu	Arg	Ser	Asp	Gly 520	Lys	Thr	Cys	Ala	Lys 525	Leu	Asp	Ser
Cys	Ala 530	Leu	Gly	Asp	His	Gly 535	Cys	Glu	His	Ser	Cys 540	Val	Ser	Ser	Glu
Asp 545	Ser	Phe	Val	Cys	Gln 550	Cys	Phe	Glu	Gly	Tyr 555	Ile	Leu	Arg	Glu	Asp 560
Gly	Lys	Thr	Cys	Arg 565	Arg	Lys	Asp	Val	Cys 570	Gln	Ala	Ile	Asp	His 575	Gly
Cys	Glu	His	Ile 580	Cys	Val	Asn	Ser	Asp 585	Asp	Ser	Tyr	Thr	Cys 590	Glu	Cys
Leu	Val	Gly 595	Phe	Arg	Leu	Ala	Glu 600	Asp	Gly	Lys	Arg	Cys 605	Arg	Arg	Lys
Asp	Val 610	Cys	Lys	Ser	Thr	His 615	His	Gly	Cys	Glu	His 620	Ile	Cys	Val	Asn
Asn 625	Gly	Asn	Ser	Tyr	Ile 630	Cys	Lys	Cys	Ser	Glu 635	Gly	Phe	Val	Leu	Ala 640

Glu	Asp	Gly	Arg	Arg 645	Cys	Lys	Lys	Cys	Thr 650	Glu	Gly	Pro	Ile	Asp 655	Leu
Val	Phe	Val	Ile 660	Asp	Gly	Ser	Lys	Ser 665	Leu	Gly	Glu	Glu	Asn 670	Phe	Glu
Val	Val	Lys 675	Gln	Phe	Val	Thr	Gly 680	Ile	Ile	Asp	Ser	Leu 685	Thr	Ile	Ser
Pro	Lys 690	Ala	Ala	Arg	Val	Gly 695	Leu	Leu	Gln	Tyr	Ser 700	Thr	Gln	Val	His
Thr 705	Glu	Phe	Thr	Leu	Arg 710	Asn	Phe	Asn	Ser	Ala 715	Lys	Asp	Met	Lys	Lys 720
Ala	Val	Ala	His	Met 725	Lys	Tyr	Met	Gly	Lys 730	Gly	Ser	Met	Thr	Gly 735	Leu
Ala	Leu	Lys	His 740	Met	Phe	Glu	Arg	Ser 745	Phe	Thr	Gln	Gly	Glu 750	Gly	Ala
Arg	Pro	Phe 755	Ser	Thr	Arg	Val	Pro 760	Arg	Ala	Ala	Ile	Val 765	Phe	Thr	Asp
Gly	Arg 770		Gln	Asp	Asp	Val 775	Ser	Glu	Trp	Ala	Ser 780	Lys	Ala	Lys	Ala
Asn 785	Gly	Ile	Thr	Met	Tyr 790	Ala	Val	Gly	Val	Gly 795	Lys	Ala	Ile	Glu	Glu 800
Glu	Leu	Gln	Glu	Ile 805	Ala	Ser	Glu	Pro	Thr 810	Asn	Lys	His	Leu	Phe 815	Tyr
Ala	Glu	Asp	Phe 820	Ser	Thr	Met	Asp	Glu 825	Ile	Ser	Glu	Lys	Leu 830	Lys	Lys
Gly	Ile	Cys 835	Glu	Ala	Leu	Glu	Asp 840	Ser	Asp	Gly	Arg	Gln 845	Asp	Ser	Pro
Ala	Gly 850	Glu	Leu	Pro	Lys	Thr 855	Val	Gln	Gln	Pro	Thr 860	Glu	Ser	Glu	Pro
Val 865	Thr	Ile	Asn	Ile	Gln 870	Asp	Leu	Leu	Ser	Cys 875	Ser	Asn	Phe	Ala	Val 880
Gln	His	Arg	Tyr	Leu 885	Phe	Glu	Glu	Asp	Asn 890	Leu	Leu	Arg	Ser	Thr 895	Gln

Lys Leu Ser His Ser Thr Lys Pro Ser Gly Ser Pro Leu Glu Glu Lys
900 905 910

His Asp Gln Cys Lys Cys Glu Asn Leu Ile Met Phe Gln Asn Leu Ala 915 920 925

Asn Glu Glu Val Arg Lys Leu Thr Gln Arg Leu Glu Glu Met Thr Gln 930 935 940

Arg Met Glu Ala Leu Glu Asn Arg Leu Arg Tyr Arg 945 950 955

<210> 75

<211> 937

<212> PRT

<213> Homo sapiens

<400> 75

Met Glu Lys Met Leu Ala Gly Cys Phe Leu Leu Ile Leu Gly Gln Ile 1 5 10 15

Val Leu Leu Pro Ala Glu Ala Arg Glu Arg Ser Arg Gly Arg Ser Ile 20 25 30

Ser Arg Gly Arg His Ala Arg Thr His Pro Gln Thr Ala Leu Leu Glu 35 40 45

Ser Ser Cys Glu Asn Lys Arg Ala Asp Leu Val Phe Ile Ile Asp Ser 50 55 60

Ser Arg Ser Val Asn Thr His Asp Tyr Ala Lys Val Lys Glu Phe Ile 65 70 75 80

Val Asp Ile Leu Gln Phe Leu Asp Ile Gly Pro Asp Val Thr Arg Val 85 90 95

Gly Leu Leu Gln Tyr Gly Ser Thr Val Lys Asn Glu Phe Ser Leu Lys 100 105 110

Thr Phe Lys Arg Lys Ser Glu Val Glu Arg Ala Val Lys Arg Met Arg 115 120 125

His Leu Ser Thr Gly Thr Met Thr Gly Leu Ala Ile Gln Tyr Ala Leu 130 135 140

Asn Ile Ala Phe Ser Glu Ala Glu Gly Ala Arg Pro Leu Arg Glu Asn 145 150 155 160

Val Pro Arg Val Ile Met Ile Val Thr Asp Gly Arg Pro Gln Asp Ser Val Ala Glu Val Ala Ala Lys Ala Arg Asp Thr Gly Ile Leu Ile Phe Ala Ile Gly Val Gly Gln Val Asp Phe Asn Thr Leu Lys Ser Ile Gly Ser Glu Pro His Glu Asp His Val Phe Leu Val Ala Asn Phe Ser Gln Ile Glu Thr Leu Thr Ser Val Phe Gln Lys Lys Leu Cys Thr Ala His Met Cys Ser Thr Leu Glu His Asn Cys Ala His Phe Cys Ile Asn Ile Pro Gly Ser Tyr Val Cys Arg Cys Lys Gln Gly Tyr Ile Leu Asn Ser Asp Gln Thr Thr Cys Arg Ile Gln Asp Leu Cys Ala Met Glu Asp His Asn Cys Glu Gln Leu Cys Val Asn Val Pro Gly Ser Phe Val Cys Gln Cys Tyr Ser Gly Tyr Ala Leu Ala Glu Asp Gly Lys Arg Cys Val Ala Val Asp Tyr Cys Ala Ser Glu Asn His Gly Cys Glu His Glu Cys Val Asn Ala Asp Gly Ser Tyr Leu Cys Gln Cys His Glu Gly Phe Ala Leu Asn Pro Asp Glu Lys Thr Cys Thr Lys Ile Asp Tyr Cys Ala Ser Ser Asn His Gly Cys Gln His Glu Cys Val Asn Thr Asp Asp Ser Tyr Ser Cys His Cys Leu Lys Gly Phe Thr Leu Asn Pro Asp Lys Lys Thr Cys Arg Arg Ile Asn Tyr Cys Ala Leu Asn Lys Pro Gly Cys Glu His Glu

Cys	Val	Asn	Met 420	Glu	Glu	Ser	Tyr	Tyr 425	Cys	Arg	Cys	His	Arg 430	Gly	Tyr
Thr	Leu	Asp 435	Pro	Asn	Gly	Lys	Thr 440	Cys	Ser	Arg	Val	Asp 445	His	Cys	Ala
Gln	Gln 450	Asp	His	Gly	Cys	Glu 455	Gln	Leu	Cys	Leu	Asn 460	Thr	Glu	Asp	Ser
Phe 465	Val	Cys	Gln	Cys	Ser 470	Glu	Gly	Phe	Leu	Ile 475	Asn	Glu	Asp	Leu	Lys 480
Thr	Cys	Ser	Arg	Val 485	Asp	Tyr	Cys	Leu	Leu 490	Ser	Asp	His	Gly	Cys 495	Glu
Tyr	Ser	Cys	Val 500	Asn	Met	Asp	Arg	Ser 505	Phe	Ala	Cys	Gln	Cys 510	Pro	Glu
Gly	His	Val 515	Leu	Arg	Ser	Asp	Gly 520	Lys	Thr	Cys	Ala	Lys 525	Leu	Asp	Ser
Cys	Ala 530	Leu	Gly	Asp	His	Gly 535	Cys	Glu	His	Ser	Cys 540	Val	Ser	Ser	Glu
Asp 545	Ser	Phe	Val	Cys	Gln 550	Cys	Phe	Glu	Gly	Tyr 555	Ile	Leu	Arg	Glu	Asp 560
Gly	Lys	Thr	Cys	Arg 565	Arg	Lys	Asp	Val	Cys 570	Gln	Ala	Ile	Asp	His 575	Gly
Cys	Glu	His	Ile 580	Cys	Val	Asn	Ser	Asp 585	Asp	Ser	Tyr	Thr	Cys 590	Glu	Cys
Leu	Glu	Gly 595	Phe	Arg	Leu	Ala	Glu 600	Asp	Gly	Lys	_	Cys 605	Arg	Arg	Lys
Asp	Val 610	Cys	Lys	Ser	Thr	His 615	His	Gly	Суз	Glu	His 620	Ile	Cys	Val	Asn
Asn 625	Gly	Asn	Ser	Tyr	Ile 630	Cys	Lys	Cys	Ser	Glu 635	Gly	Phe	Val	Leu	Ala 640
Glu	Asp	Gly	Arg	Arg 645	Cys	Lys	Lys	Cys	Thr 650	Glu	Gly	Pro	Ile	Asp 655	Leu
Val	Phe	Val	Ile 660	Asp	Gly	Ser	Lys	Ser 665	Leu	Gly	Glu	Glu	Asn 670	Phe	Glu

Val Val Lys Gln Phe Val Thr Gly Ile Ile Asp Ser Leu Thr Ile Ser Pro Lys Ala Ala Arq Val Gly Leu Leu Gln Tyr Ser Thr Gln Val His Thr Glu Phe Thr Leu Arg Asn Phe Asn Ser Ala Lys Asp Met Lys Lys Ala Val Ala His Met Lys Tyr Met Gly Lys Gly Ser Met Thr Gly Leu Ala Leu Lys His Met Phe Glu Arg Ser Phe Thr Gln Gly Glu Gly Ala Arg Pro Leu Ser Thr Arg Val Pro Arg Ala Ala Ile Val Phe Thr Asp Gly Arg Ala Gln Asp Asp Val Ser Glu Trp Ala Ser Lys Ala Lys Ala Asn Gly Ile Thr Met Tyr Ala Val Gly Val Gly Lys Ala Ile Glu Glu Glu Leu Gln Glu Ile Ala Ser Glu Pro Thr Asn Lys His Leu Phe Tyr Ala Glu Asp Phe Ser Thr Met Asp Glu Ile Ser Glu Lys Leu Lys Lys Gly Ile Cys Glu Ala Leu Glu Asp Ser Asp Gly Arg Gln Asp Ser Pro Ala Gly Glu Leu Pro Lys Thr Val Gln Pro Thr Val Gln His Arg Tyr Leu Phe Glu Glu Asp Asn Leu Leu Arg Ser Thr Gln Lys Leu Ser His Ser Thr Lys Pro Ser Gly Ser Pro Leu Glu Glu Lys His Asp Gln Cys Lys Cys Glu Asn Leu Ile Met Phe Gln Asn Leu Ala Asn Glu Glu Val Arg Lys Leu Thr Gln Arg Leu Glu Glu Met Thr Gln Arg Met Glu

Ala Leu Glu Asn Arg Leu Arg Tyr Arg 930 935

<210> 76

<211> 956

<212> PRT

<213> Mus musculus

<400> 76

Met Glu Lys Met Leu Val Gly Cys Leu Leu Met Leu Gly Gln Leu Phe 1 5 10 15

Leu Val Leu Pro Val Asp Gly Arg Glu Arg Pro Gln Ala Arg Phe Pro 20 25 30

Ser Arg Gly Arg His Val Arg Met Tyr Pro Gln Thr Ala Leu Leu Glu 35 40 45

Ser Ser Cys Glu Asn Lys Arg Ala Asp Leu Val Phe Ile Ile Asp Ser 50 55 60

Ser Arg Ser Val Asn Thr His Asp Tyr Ala Lys Val Lys Glu Phe Ile 65 70 75 80

Leu Asp Ile Leu Gln Phe Leu Asp Ile Gly Pro Asp Val Thr Arg Val
85 90 95

Gly Leu Leu Gln Tyr Gly Ser Thr Val Lys Asn Glu Phe Ser Leu Lys 100 105 110

Thr Phe Lys Arg Lys Ser Glu Val Glu Arg Ala Val Lys Arg Met Arg
115 120 125

His Leu Ser Thr Gly Thr Met Thr Gly Leu Ala Ile Gln Tyr Ala Leu 130 135 140

Val Pro Arg Ile Ile Met Ile Val Thr Asp Gly Arg Pro Gln Asp Ser 165 170 175

Val Ala Glu Val Ala Ala Lys Ala Arg Asn Thr Gly Ile Leu Ile Phe 180 185 190

Ala Ile Gly Val Gly Gln Val Asp Leu Asn Thr Leu Lys Ala Ile Gly

		195					200					205			
Ser	Glu 210	Pro	His	Lys	Asp	His 215	Val	Phe	Leu	Val	Ala 220	Asn	Phe	Ser	Gln
Ile 225	Glu	Ser	Leu	Thr	Ser 230	Val	Phe	Gln	Asn	Lys 235	Leu	Cys	Thr	Val	His 240
Met	Cys	Ser	Ile	Leu 245	Glu	His	Asn	Cys	Ala 250	His	Phe	Cys	Leu	Asn 255	Thr
Pro	Gly	Ser	Tyr 260	Ile	Cys	Lys	Cys	Lys 265	Gln	Gly	Tyr	Ile	Leu 270	Ser	Thr
Asp	Gln	Lys 275	Thr	Cys	Arg	Ile	Gln 280	Asp	Leu	Cys	Ala	Thr 285	Glu	Asp	His
Gly	Cys 290	Glu	Gln	Leu	Cys	Val 295	Asn	Met	Leu	Gly	Ser 300	Phe	Val	Cys	Gln
Cys 305	Tyr	Ser	Gly	Tyr	Thr 310	Leu	Ala	Glu	Asp	Gly 315	Lys	Arg	Cys	Thr	Ala 320
Val	Asp	Tyr	Cys	Ala 325	Ser	Glu	Asn	His	Gly 330	Cys	Glu	His	Glu	Cys 335	Val
Asn	Ala	Glu	Ser 340	Ser	Tyr	Leu	Cys	Arg 345	Cys	His	Glu	Gly	Phe 350	Ala	Leu
Asn	Ser	Asp 355	Lys	Lys	Thr	Cys	Ser 360	Lys	Ile	Asp	Tyr	Cys 365	Ala	Ser	Ser
Asn	His 370	Gly	Cys	Gln	His	Glu 375	Cys	Val	Asn	Ala	Gln 380	Thr	Ser	Ala	Leu
Cys 385	Arg	Cys	Leu	Lys	Gly 390	Phe	Met	Leu	Asn	Pro 395	Asp	Arg	Lys	Thr	Cys 400
Arg	Arg	Ile	Asn	Tyr 405	Cys	Ala	Leu	Asn	Lys 410	Pro	Gly	Cys	Glu	His 415	Glu
Cys	Val	Asn	Thr 420	Glu	Glu	Gly	His	Tyr 425	Cys	Arg	Cys	Arg	Gln 430	Gly	Tyr
Asn	Leu	Asp 435	Pro	Asn	Gly	Lys	Thr 440	Cys	Ser	Arg	Val	Asp 445	His	Cys	Ala

Gln Gln Asp His Gly Cys Glu Gln Leu Cys Leu Asn Thr Glu Glu Ser

Phe 465	Val	Cys	Gln	Суз	Ser 470	Glu	Gly	Phe	Leu	Ile 475	Asn	Asp	Asp	Leu	Lys 480
Thr	Cys	Ser	Arg	Ala 485	Asp	Tyr	Cys	Leu	Leu 490	Ser	Asn	His	Gly	Cys 495	Glu
Tyr	Ser	Cys	Val 500	Asn	Thr	Asp	Lys	Ser 505	Phe	Ala	Cys	Gln	Cys 510	Pro	Glu
Gly	His	Val 515	Leu	Arg	Ser	Asp	Gly 520	Lys	Thr	Cys	Ala	Lys 525	Leu	Asp	Ser
Cys	Ala 530	Leu	Gly	Asp	His	Gly 535	Cys	Glu	His	Ser	Cys 540	Val	Ser	Ser	Glu
Asp 545	Ser	Phe	Val	Cys	Gln 550	Cys	Phe	Glu	Gly	Tyr 555	Ile	Leu	Arg	Asp	Asp 560
Gly	Lys	Thr	Cys	Arg 565	Arg	Lys	Asp	Val	Cys 570	Gln	Asp	Val	Asn	His 575	Gly
Cys	Glu	His	Leu 580	Cys	Val	Asn	Ser	Gly 585	Glu	Ser	Tyr	Val	Cys 590	Lys	Cys
Leu	Glu	Gly 595	Phe	Arg	Leu	Ala	Glu 600	Asp	Gly	Lys	Arg	Cys 605	Arg	Arg	Lys
Asn	Val 610	Cys	Lys	Ser	Thr	Gln 615	His	Gly	Cys	Glu	His 620	Met	Cys	Val	Asn
Asn 625	Gly	Asn	Ser	Tyr	Leu 630	Cys	Arg	Cys	Ser	Glu 635	Gly	Phe	Val	Leu	Ala 640
Glu	Asp	Gly	Lys	His 645	Cys	Lys	Arg	Cys	Thr 650	Glu	Gly	Pro	Ile	Asp 655	Leu
Val	Phe	Val	Ile 660	Asp	Gly	Ser	Lys	Ser 665	Leu	Gly	Glu	Glu	Asn 670	Phe	Glu
Thr	Val	Lys 675	His	Phe	Val	Thr	Gly 680	Ile	Ile	Asp	Ser	Leu 685	Ala	Val	Ser
Pro	Lys 690	Ala	Ala	Arg	Val	Gly 695	Leu	Leu	Gln	Tyr	Ser 700	Thr	Gln	Val	Arg
Thr	Glu	Phe	Thr	Leu	Arg	Gly	Phe	Ser	Ser	Ala	Lys	Glu	Met	Lys	Lys

705		710				715					720
Ala Val <i>F</i>	Ala His	Met Lys 725	Tyr Me	t Gly	Lys 730	Gly	Ser	Met	Thr	Gly 735	Leu
Ala Leu I	Lys His 740	Met Phe	Glu Ar	g Ser 745	Phe	Thr	Gln	Val	Glu 750	Gly	Ala
Arg Pro I	Pro Ser 755	Thr Gln	Val Pr 76	_	Val	Ala	Ile	Val 765	Phe	Thr	Asp
Gly Arg 770	Ala Gln	Asp Asp	Val Se 775	r Glu	Trp	Ala	Ser 780	Lys	Ala	Lys	Ala
Asn Gly 1 785	Ile Thr	Met Tyr 790	Ala Va	l Gly	Val	Gly 795	Lys	Ala	Ile	Glu	Glu 800
Glu Leu G	Gln Glu	Ile Ala 805	Ser Gl	u Pro	Ile 810	Asp	Lys	His	Leu	Phe 815	Tyr
Ala Glu <i>F</i>	Asp Phe 820	Ser Thr	Met Gl	y Glu 825	Ile	Ser	Glu	Lys	Leu 830	Lys	Glu
Gly Ile C	Cys Glu 335	Ala Leu	Glu As 84		Gly	Gly	Arg	Gln 845	Asp	Ser	Ala
Ala Trp A 850	Asp Leu	Pro Gln	Gln Al 855	a His	Gln	Pro	Thr 860	Glu	Pro	Glu	Pro
Val Thr I 865	Ile Lys	Ile Lys 870	Asp Le	u Leu	Ser	Cys 875	Ser	Asn	Phe	Ala	Val 880
Gln His A	_	Leu Phe 885	Glu Gl	u Asp	Asn 890	Leu	Ser	Arg	Ser	Thr 895	Gln
Lys Leu F	Phe His	Ser Thr	Lys Se	r Ser 905	Gly	Asn	Pro	Leu	Glu 910	Glu	Ser
Gln Asp G	Gln Cys 915	Lys Cys	Glu As 92		Ile	Leu	Phe	Gln 925	Asn	Val	Ala
Asn Glu G 930	Glu Val	Arg Lys	Leu Th	r Gln	Arg	Leu	Glu 940	Glu	Met	Thr	Gln
Arg Met G	Glu Ala	Leu Glu 950	Asn Ar	g Leu	Lys	Tyr 955	Arg				

<210> 77

<211> 956

<212> PRT

<213> Mus musculus

<400> 77

Met Glu Lys Met Leu Val Gly Cys Leu Leu Met Leu Gly Gln Leu Phe $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Leu Val Leu Pro Val Asp Gly Arg Glu Arg Pro Gln Ala Arg Phe Pro 20 25 30

Ser Arg Gly Arg His Val Arg Met Tyr Pro Gln Thr Ala Leu Leu Glu 35 40 45

Ser Ser Cys Glu Asn Lys Arg Ala Asp Leu Val Phe Ile Ile Asp Ser 50 55 60

Ser Arg Ser Val Asn Thr Tyr Asp Tyr Ala Lys Val Lys Glu Phe Ile 65 70 75 80

Leu Asp Ile Leu Gln Phe Leu Asp Ile Gly Pro Asp Val Thr Arg Val 85 90 95

Gly Leu Leu Gln Tyr Gly Ser Thr Val Lys Asn Glu Phe Ser Leu Lys 100 105 110

Thr Phe Lys Arg Lys Ser Glu Val Glu Arg Ala Val Lys Arg Met Arg 115 120 125

His Leu Ser Thr Gly Thr Met Thr Gly Leu Ala Ile Gln Tyr Ala Leu 130 135 140

Asn Ile Ala Phe Ser Glu Ala Glu Gly Ala Arg Pro Leu Arg Glu Asn 145 150 155 160

Val Pro Arg Ile Ile Met Ile Val Thr Asp Gly Arg Pro Gln Asp Ser 165 170 · 175

Val Ala Glu Val Ala Ala Lys Ala Arg Asn Thr Gly Ile Leu Ile Phe 180 185 190

Ala Ile Gly Val Gly Gln Val Asp Leu Asn Thr Leu Lys Ala Ile Gly 195 200 205

Ser Glu Pro His Lys Asp His Val Phe Leu Val Ala Asn Phe Ser Gln 210 215 220

Ile 225	Glu	Ser	Leu	Thr	Ser 230	Val	Phe	Gln	Asn	Lys 235	Leu	Cys	Thr	Val	His 240
Met	Cys	Ser	Val	Leu 245	Glu	His	Asn	Cys	Ala 250	His	Phe	Cys	Leu	Asn 255	Thr
Pro	Gly	Ser	Tyr 260	Ile	Cys	Lys	Суз	Lys 265	Gln	Gly	Tyr	Ile	Leu 270	Ser	Thr
Asp	Gln	Lys 275	Thr	Cys	Arg	Ile	Gln 280	Asp	Leu	Cys	Ala	Thr 285	Glu	Asp	His
Gly	Cys 290	Glu	Gln	Leu	Cys	Val 295	Asn	Met	Leu	Gly	Ser 300	Phe	Val	Cys	Gln
Cys 305	Tyr	Ser	Gly	Tyr	Thr 310	Leu	Ala	Glu	Asp	Gly 315	Lys	Arg	Cys	Thr	Ala 320
Met	Asp	Tyr	Cys	Ala 325	Ser	Glu	Asn	His	Gly 330	Cys	Glu	His	Glu	Cys 335	Val
Asn	Ala	Glu	Ser 340	Ser	Tyr	Leu	Cys	Arg 345	Cys	His	Glu	Gly	Phe 350	Ala	Leu
Asn	Ser	Asp 355	Lys	Lys	Thr	Cys	Ser 360	Lys	Ile	Asp	Tyr	Cys 365	Ala	Ser	Ser
Asn	His 370	Gly	Cys	Gln	His	Glu 375	Cys	Val	Asn	Ala	Gln 380	Thr	Ser	Ala	Leu
Cys 385	Arg	Cys	Leu	Lys	Gly 390	Phe	Met	Leu	Asn	Pro 395	Asp	Arg	Lys	Thr	Cys 400
Arg	Arg	Ile	Asn	Tyr 405	Суѕ	Ala	Leu	Asn	Lys 410	Pro	Gly	Cys	Glu	His 415	Glu
Cys	Val	Asn	Thr 420	Glu	Glu	Gly	His	Tyr 425	Cys	Arg	Cys	Arg	Gln 430	Gly	Tyr
Asn	Leu	Asp 435	Pro	Asn	Gly	Lys	Thr 440	Cys	Ser	Arg	Val	Asp 445	His	Cys	Ala
Gln	Gln 450	Asp	His	Gly	Cys	Glu 455	Gln	Leu	Cys	Leu	Asn 460	Thr	Glu	Glu	Ser
Phe 465	Val	Cys	Gln	Cys	Ser 470	Glu	Gly	Phe	Leu	Ile 475	Asn	Asp	Asp	Leu	Lys 480

Thr Cys Ser Arg Ala Asp Tyr Cys Leu Leu Ser Asn His Gly Cys Glu Tyr Ser Cys Val Asn Thr Asp Lys Ser Phe Ala Cys Gln Cys Pro Glu Gly His Val Leu Arg Ser Asp Gly Lys Thr Cys Ala Lys Leu Asp Ser Cys Ala Leu Gly Asp His Gly Cys Glu His Ser Cys Val Ser Ser Glu Asp Ser Phe Val Cys Gln Cys Phe Glu Gly Tyr Ile Leu Arg Asp Asp Gly Lys Thr Cys Arg Arg Lys Asp Val Cys Gln Asp Val Asn His Gly Cys Glu His Leu Cys Val Asn Ser Gly Glu Ser Tyr Val Cys Lys Cys Leu Glu Gly Phe Arg Leu Ala Glu Asp Gly Lys Arg Cys Arg Arg Lys Asn Val Cys Lys Ser Thr Gln His Gly Cys Glu His Met Cys Val Asn Asn Gly Asn Ser Tyr Leu Cys Arg Cys Ser Glu Gly Phe Val Leu Ala Glu Asp Gly Lys His Cys Lys Arg Cys Thr Glu Gly Pro Ile Asp Leu Val Phe Val Ile Asp Gly Ser Lys Ser Leu Gly Glu Glu Asn Phe Glu Thr Val Lys His Phe Val Thr Gly Ile Ile Asp Ser Leu Ala Val Ser Pro Lys Ala Ala Arg Val Gly Leu Leu Gln Tyr Ser Thr Gln Val Arg Thr Glu Phe Thr Leu Arg Gly Phe Ser Ser Ala Lys Glu Met Lys Lys Ala Val Thr His Met Lys Tyr Met Gly Lys Gly Ser Met Thr Gly Leu

Ala Leu Lys His Met Phe Glu Arg Ser Phe Thr Gln Val Glu Gly Ala 740 745 750

Arg Pro Pro Ser Thr Gln Val Pro Arg Val Ala Ile Val Phe Thr Asp 755 760 765

Gly Arg Ala Gln Asp Asp Val Ser Glu Trp Ala Ser Lys Ala Lys Ala 770 780

Asn Gly Ile Thr Met Tyr Ala Val Gly Val Gly Lys Ala Ile Glu Glu 785 790 795 800

Glu Leu Gln Glu Ile Ala Ser Glu Pro Ile Asp Lys His Leu Phe Tyr 805 810 815

Ala Glu Asp Phe Ser Thr Met Gly Glu Ile Ser Glu Lys Leu Lys Glu 820 825 830

Gly Ile Cys Glu Ala Leu Glu Asp Ser Gly Gly Arg Gln Asp Ser Ala 835 840 845

Ala Trp Asp Leu Pro Gln Gln Ala His Gln Pro Thr Glu Pro Glu Pro 850 855 860

Val Thr Ile Lys Ile Lys Asp Leu Leu Ser Cys Ser Asn Phe Ala Val 865 870 875 880

Gln His Arg Phe Leu Phe Glu Glu Asp Asn Leu Ser Arg Ser Thr Gln 885 890 895

Lys Leu Phe His Ser Thr Lys Ser Ser Gly Asn Pro Leu Glu Glu Ser 900 905 910

Gln Asp Gln Cys Lys Cys Glu Asn Leu Ile Leu Phe Gln Asn Val Ala 915 920 925

Asn Glu Glu Val Arg Lys Leu Thr Gln Arg Leu Glu Glu Met Thr Gln 930 935 940

Arg Met Glu Ala Leu Glu Asn Arg Leu Lys Tyr Arg 945 950 955

<210> 78

<211> 200

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Von Willebrand Factor type A doman sequence

<400> 78

Asp Ile Val Phe Leu Leu Asp Gly Ser Gly Ser Ile Gly Ser Gln Asn 1 5 10 15

Phe Glu Arg Val Lys Asp Phe Val Glu Arg Val Val Glu Arg Leu Asp 20 25 ' 30

Val Gly Pro Arg Asp Lys Lys Glu Glu Asp Ala Val Arg Val Gly Leu 35 40 45

Val Gln Tyr Ser Asp Asn Val Arg Thr Glu Ile Lys Phe Lys Leu Asn 50 55 60

Asp Tyr Gln Asn Lys Asp Glu Val Leu Gln Ala Leu Gln Lys Ile Arg
65 70 75 80

Tyr Glu Asp Tyr Tyr Gly Gly Gly Gly Thr Asn Thr Gly Ala Ala Leu 85 90 95

Gln Tyr Val Val Arg Asn Leu Phe Thr Glu Ala Ser Gly Ser Arg Ile 100 105 110

Glu Pro Val Ala Glu Glu Gly Ala Pro Lys Val Leu Val Val Leu Thr
115 120 125

Asp Gly Arg Ser Gln Asp Asp Pro Ser Pro Thr Ile Asp Ile Arg Asp 130 135 140

Val Leu Asn Glu Leu Lys Lys Glu Ala Gly Val Glu Val Phe Ala Ile 145 150 155 160

Gly Val Gly Asn Ala Asp Asn Asn Asn Leu Glu Glu Leu Arg Glu Ile 165 170 175

Ala Ser Lys Pro Asp Asp His Val Phe Lys Val Ser Asp Phe Glu Ala 180 185 190

Leu Asp Thr Leu Gln Glu Leu Leu 195 200

<210> 79

<211> 176

<212> PRT

<213> Homo sapiens

<400> 79

Asp Leu Val Phe Ile Ile Asp Ser Ser Arg Ser Val Asn Thr His Asp 1 5 10 15

Tyr Ala Lys Val Lys Glu Phe Ile Val Asp Ile Leu Gln Phe Leu Asp 20 25 30

Ile Gly Pro Asp Val Thr Arg Val Gly Leu Leu Gln Tyr Gly Ser Thr 35 40 45

Val Lys Asn Glu Phe Ser Leu Lys Thr Phe Lys Arg Lys Ser Glu Val
50 55 60

Glu Arg Ala Val Lys Arg Met Arg His Leu Ser Thr Gly Thr Met Thr
65 70 75 80

Gly Leu Ala Ile Gln Tyr Ala Leu Asn Ile Ala Phe Ser Glu Ala Glu 85 90 95.

Gly Ala Arg Pro Leu Arg Glu Asn Val Pro Arg Val Ile Met Ile Val 100 \$100\$

Thr Asp Gly Arg Pro Gln Asp Ser Val Ala Glu Val Ala Ala Lys Ala 115 120 125

Arg Asp Thr Gly Ile Leu Ile Phe Ala Ile Gly Val Gly Gln Val Asp 130 135 140

Phe Asn Thr Leu Lys Ser Ile Gly Ser Glu Pro His Glu Asp His Val 145 150 155 160

Phe Leu Val Ala Asn Phe Ser Gln Ile Glu Thr Leu Thr Ser Val Phe 165 170 175

<210> 80

<211> 200

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Von Willebrand Factor type A doman sequence

<400> 80

Asp Ile Val Phe Leu Leu Asp Gly Ser Gly Ser Ile Gly Ser Gln Asn 1 5 10 15

Phe Glu Arg Val Lys Asp Phe Val Glu Arg Val Glu Arg Leu Asp 20 25 30

Val Gly Pro Arg Asp Lys Lys Glu Glu Asp Ala Val Arg Val Gly Leu 35 40 45

Val Gln Tyr Ser Asp Asn Val Arg Thr Glu Ile Lys Phe Lys Leu Asn 50 55 60

Asp Tyr Gln Asn Lys Asp Glu Val Leu Gln Ala Leu Gln Lys Ile Arg 65 70 75 80

Tyr Glu Asp Tyr Tyr Gly Gly Gly Gly Thr Asn Thr Gly Ala Ala Leu 85 90 95

Gln Tyr Val Val Arg Asn Leu Phe Thr Glu Ala Ser Gly Ser Arg Ile 100 105 110

Glu Pro Val Ala Glu Glu Gly Ala Pro Lys Val Leu Val Val Leu Thr
115 120 125

Asp Gly Arg Ser Gln Asp Asp Pro Ser Pro Thr Ile Asp Ile Arg Asp 130 135 140

Val Leu Asn Glu Leu Lys Lys Glu Ala Gly Val Glu Val Phe Ala Ile 145 150 155 160

Gly Val Gly Asn Ala Asp Asn Asn Leu Glu Glu Leu Arg Glu Ile 165 170 175

Ala Ser Lys Pro Asp Asp His Val Phe Lys Val Ser Asp Phe Glu Ala 180 185 190

Leu Asp Thr Leu Gln Glu Leu Leu 195 200

<210> 81

<211> 176

<212> PRT

<213> Homo sapiens

<400> 81

Asp Leu Val Phe Val Ile Asp Gly Ser Lys Ser Leu Gly Glu Glu Asn 1 5 10 15

Phe Glu Val Val Lys Gln Phe Val Thr Gly Ile Ile Asp Ser Leu Thr 20 25 30

Ile Ser Pro Lys Ala Ala Arg Val Gly Leu Leu Gln Tyr Ser Thr Gln
35 40 45

Val His Thr Glu Phe Thr Leu Arg Asn Phe Asn Ser Ala Lys Asp Met 50 55 60

Lys Lys Ala Val Ala His Met Lys Tyr Met Gly Lys Gly Ser Met Thr 65 70 75 80

Gly Leu Ala Leu Lys His Met Phe Glu Arg Ser Phe Thr Gln Gly Glu 85 90 95

Gly Ala Arg Pro Phe Ser Thr Arg Val Pro Arg Ala Ala Ile Val Phe 100 105 110

Thr Asp Gly Arg Ala Gln Asp Asp Val Ser Glu Trp Ala Ser Lys Ala 115 120 125

Lys Ala Asn Gly Ile Thr Met Tyr Ala Val Gly Val Gly Lys Ala Ile 130 135 140

Phe Tyr Ala Glu Asp Phe Ser Thr Met Asp Glu Ile Ser Glu Lys Leu 165 170 175

<210> 82

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: EGF domain
 sequence

<400> 82

Cys Ala Pro Asn Asn Pro Cys Ser Asn Gly Gly Thr Cys Val Asn Thr

1 5 10 15

Pro Gly Gly Ser Ser Asp Asn Phe Gly Gly Tyr Thr Cys Glu Cys Pro
20 25 30

Pro Gly Asp Tyr Tyr Leu Ser Tyr Thr Gly Lys Arg Cys 35 40 45

<210> 83

<211> 36

<212> PRT

<213> Homo sapiens

<400> 83

Cys Ser Thr Leu Glu His Asn Cys Ala His Phe Cys Ile Asn Ile Pro 1 5 10 15

Gly Ser Tyr Val Cys Arg Cys Lys Gln Gly Tyr Ile Leu Asn Ser Asp 20 25 30

Gln Thr Thr Cys 35

<210> 84

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: EGF domain sequence

<400> 84

Cys Ala Pro Asn Asn Pro Cys Ser Asn Gly Gly Thr Cys Val Asn Thr
1 5 10 15

Pro Gly Gly Ser Ser Asp Asn Phe Gly Gly Tyr Thr Cys Glu Cys Pro
20 25 30

Pro Gly Asp Tyr Tyr Leu Ser Tyr Thr Gly Lys Arg Cys
35 40 45

<210> 85

<211> 36

<212> PRT

<213> Homo sapiens <400> 85 Cys Ala Met Glu Asp His Asn Cys Glu Gln Leu Cys Val Asn Val Pro 5 10 Gly Ser Phe Val Cys Gln Cys Tyr Ser Gly Tyr Ala Leu Ala Glu Asp Gly Lys Arg Cys 35 <210> 86 <211> 45 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: EGF domain sequence <400> 86 Cys Ala Pro Asn Asn Pro Cys Ser Asn Gly Gly Thr Cys Val Asn Thr 5 Pro Gly Gly Ser Ser Asp Asn Phe Gly Gly Tyr Thr Cys Glu Cys Pro 20 30 Pro Gly Asp Tyr Tyr Leu Ser Tyr Thr Gly Lys Arg Cys 35 40 <210> 87 <211> 36 <212> PRT <213> Homo sapiens Cys Ala Ser Glu Asn His Gly Cys Glu His Glu Cys Val Asn Ala Asp 1 15 Gly Ser Tyr Leu Cys Gln Cys His Glu Gly Phe Ala Leu Asn Pro Asp

25

131

30

20

Glu Lys Thr Cys 35 <210> 88

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: EGF domain sequence

<400> 88

Cys Ala Pro Asn Asn Pro Cys Ser Asn Gly Gly Thr Cys Val Asn Thr 1 5 10 15

Pro Gly Gly Ser Ser Asp Asn Phe Gly Gly Tyr Thr Cys Glu Cys Pro 20 25 30

Pro Gly Asp Tyr Tyr Leu Ser Tyr Thr Gly Lys Arg Cys 35 40 45

<210> 89

<211> 36

<212> PRT

<213> Homo sapiens

<400> 89

Cys Ala Ser Ser Asn His Gly Cys Gln Tyr Glu Cys Val Asn Thr Asp $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asp Ser Tyr Ser Cys His Cys Leu Lys Gly Phe Thr Leu Asn Pro Asp 20 25 30

Lys Lys Thr Cys 35

<210> 90

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: EGF domain sequence

<400> 90

Cys Ala Pro Asn Asn Pro Cys Ser Asn Gly Gly Thr Cys Val Asn Thr

1 5 10 15

Pro Gly Gly Ser Ser Asp Asn Phe Gly Gly Tyr Thr Cys Glu Cys Pro
20 25 30

Pro Gly Asp Tyr Tyr Leu Ser Tyr Thr Gly Lys Arg Cys
35 40 45

<210> 91

<211> 81

<212> PRT

<213> Homo sapiens

<400> 91

Cys Ala Leu Asn Lys Pro Gly Cys Glu Cys Ala Pro Asn Asn Pro Cys
1 5 10 15

Ser Asn Gly Gly Thr Cys Val Asn Thr Pro Gly Gly Ser Ser Asp Asn 20 25 30

Phe Gly Gly Tyr Thr Cys Glu Cys Pro Pro Gly Asp Tyr Tyr Leu Ser
35 40 45

Tyr Thr Gly Lys Arg Cys His Glu Cys Val Asn Met Glu Glu Ser Tyr 50 55 60

Tyr Cys Arg Cys His Arg Gly Tyr Thr Leu Asp Pro Asn Gly Lys Pro 65 70 75 80

Cys

<210> 92

<211> 81

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: EGF domain sequence

<400> 92

Cys Ala Leu Asn Lys Pro Gly Cys Glu Cys Ala Pro Asn Asn Pro Cys
1 5 10 15

Ser Asn Gly Gly Thr Cys Val Asn Thr Pro Gly Gly Ser Ser Asp Asn

20

25

30

Phe Gly Gly Tyr Thr Cys Glu Cys Pro Pro Gly Asp Tyr Tyr Leu Ser 35 40 45

Tyr Thr Gly Lys Arg Cys His Glu Cys Val Asn Met Glu Glu Ser Tyr 50 55 60

Tyr Cys Arg Cys His Arg Gly Tyr Thr Leu Asp Pro Asn Gly Lys Pro 65 70 75 80

Cys

<210> 93

<211> 36

<212> PRT

<213> Homo sapiens

<400> 93

Cys Ala Gln Gln Asp His Gly Cys Glu Gln Leu Cys Leu Asn Thr Glu
1 5 10 15

Asp Ser Phe Val Cys Gln Cys Ser Glu Gly Phe Leu Ile Asn Glu Asp 20 25 30

Leu Lys Thr Cys 35

<210> 94

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: EGF domain
 sequence

<400> 94

Cys Ala Pro Asn Asn Pro Cys Ser Asn Gly Gly Thr Cys Val Asn Thr
1 5 10 15

Pro Gly Gly Ser Ser Asp Asn Phe Gly Gly Tyr Thr Cys Glu Cys Pro 20 25 30

Pro Gly Asp Tyr Tyr Leu Ser Tyr Thr Gly Lys Arg Cys

35

40

45

<210> 95

<211> 36

<212> PRT

<213> Homo sapiens

<400> 95

Cys Leu Leu Ser Asp His Gly Cys Glu Tyr Ser Cys Val Asn Met Asp 1 5 10 15

Arg Ser Phe Ala Cys Gln Cys Pro Glu Gly His Val Leu Arg Ser Asp 20 25 30

Gly Lys Thr Cys 35

<210> 96

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: EGF domain
 sequence

<400> 96

Cys Ala Pro Asn Asn Pro Cys Ser Asn Gly Gly Thr Cys Val Asn Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Pro Gly Gly Ser Ser Asp Asn Phe Gly Gly Tyr Thr Cys Glu Cys Pro
20 25 30

Pro Gly Asp Tyr Tyr Leu Ser Tyr Thr Gly Lys Arg Cys 35 40 45

<210> 97

<211> 36

<212> PRT

<213> Homo sapiens

<400> 97

Cys Ala Leu Gly Asp His Gly Cys Glu His Ser Cys Val Ser Ser Glu
1 5 10 15

Asp Ser Phe Val Cys Gln Cys Phe Glu Gly Tyr Ile Leu Arg Glu Asp 20 25 30

Gly Lys Thr Cys 35

<210> 98

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: EGF domain sequence

<400> 98

Cys Ala Pro Asn Asn Pro Cys Ser Asn Gly Gly Thr Cys Val Asn Thr
1 5 10 15

Pro Gly Gly Ser Ser Asp Asn Phe Gly Gly Tyr Thr Cys Glu Cys Pro 20 25 30

Pro Gly Asp Tyr Tyr Leu Ser Tyr Thr Gly Lys Arg Cys 35 40 45

<210> 99

<211> 36

<212> PRT

<213> Homo sapiens

<400> 99

Cys Gln Ala Ile Asp His Gly Cys Glu His Ile Cys Val Asn Ser Asp 1 5 10 15

Asp Ser Tyr Thr Cys Glu Cys Leu Glu Gly Phe Arg Leu Thr Glu Asp 20 25 30

Gly Lys Arg Cys 35

<210> 100

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: EGF domain sequence

<400> 100

Cys Ala Pro Asn Asn Pro Cys Ser Asn Gly Gly Thr Cys Val Asn Thr
1 5 10 15

Pro Gly Gly Ser Ser Asp Asn Phe Gly Gly Tyr Thr Cys Glu Cys Pro
20 25 30

Pro Gly Asp Tyr Tyr Leu Ser Tyr Thr Gly Lys Arg Cys 35 40 45

<210> 101

<212> PRT

<213> Homo sapiens

<400> 101

<210> 102

<211> 464

<212> PRT

<213> Rattus norvegicus

<400> 102

Met Val Leu Ala Phe Trp Leu Ala Phe Phe Thr Tyr Thr Trp Ile Thr 1 5 10 15

Leu Met Leu Asp Ala Ser Ala Val Lys Glu Pro His Gln Gln Cys Leu 20 25 30

Ser Ser Pro Lys Gln Thr Arg Ile Arg Glu Thr Arg Met Arg Lys Asp 35 40 45

Asp Leu Thr Lys Val Trp Pro Leu Lys Arg Glu Gln Leu Leu His Ile 50 55 60

Glu Asp His Asp Phe Ser Thr Arg Pro Gly Phe Gly Gly Ser Pro Val 65 70 75 80

Pro Val Gly Ile Asp Val Gln Val Glu Ser Ile Asp Ser Ile Ser Glu 85 90 95

Val Asn Met Asp Phe Thr Met Thr Phe Tyr Leu Arg His Tyr Trp Lys

			100					103					110		
Asp	Glu	Arg 115	Leu	Ser	Phe	Pro	Ser 120	Thr	Thr	Asn	Lys	Ser 125	Met	Thr	Phe
Asp	Arg 130	Arg	Leu	Ile	Gln	Lys 135	Ile	Trp	Val	Pro	Asp 140	Ile	Phe	Phe	Val
His 145	Ser	Lys	Arg	Ser	Phe 150	Ile	His	Asp	Thr	Thr 155	Val	Glu	Asn	Ile	Met 160
Leu	Arg	Val.	His	Pro 165	Asp	Gly	Asn	Val	Leu 170	Phe	Ser	Leu	Arg	Ile 175	Thr
Val	Ser	Ala	Met 180	Cys	Phe	Met	Asp	Phe 185	Ser	Arg	Phe	Pro	Leu 190	Asp	Thr
Gln	Asn	Cys 195	Ser	Leu	Glu	Leu	Glu 200	Ser	Tyr	Ala	Tyr	Asn 205	Glu	Glu	Asp
Leu	Met 210	Leu	Tyr	Trp	Lys	His 215	Gly	Asn	Lys	Ser	Leu 220	Asn	Thr	Glu	Glu
His 225	Ile	Ser	Leu	Ser	Gln 230	Phe	Phe	Ile	Glu	Glu 235	Phe	Ser	Ala	Ser	Ser 240
Gly	Leu	Ala	Phe	Tyr 245	Ser	Ser	Thr	Gly	Trp 250	Tyr	Tyr	Arg	Leu	Phe 255	Ile
Asn	Phe	Val	Leu 260	Arg	Arg	His	Ile	Phe 265	Phe	Phe	Val	Leu	Gln 270	Thr	Tyr
Phe	Pro	Ala 275	Met	Leu	Met	Val	Met 280	Leu	Ser	Trp	Val	Ser 285	Phe	Trp	Ile
Asp	Arg 290	Arg	Ala	Val	Pro	Ala 295	Arg	Val	Ser	Leu	Gly 300	Ile	Thr	Thr	Val
Leu 305	Thr	Met	Ser	Thr	Ile 310	Val	Thr	Gly	Val	Ser 315	Ala	Ser	Met	Pro	Gln 320
Val	Ser	Tyr	Val	Lys 325	Ala	Val	Asp	Val	Tyr 330	Met	Trp	Val	Ser	Ser 335	Leu

Phe Val Phe Leu Ser Val Ile Glu Tyr Ala Ala Val Asn Tyr Leu Thr

Thr Val Glu Glu Trp Lys Gln Leu Asn Arg Arg Gly Lys Ile Ser Gly

355 360 365

Met Tyr Asn Ile Asp Ala Val Gln Ala Met Ala Phe Asp Gly Cys Tyr 370 375 380

His Asp Gly Glu Thr Asp Val Asp Gln Thr Ser Phe Phe Leu His Ser 385 390 395 400

Glu Glu Asp Ser Met Arg Thr Lys Phe Thr Gly Ser Pro Cys Ala Asp
405 410 415

Ser Ser Gln Ile Lys Arg Lys Ser Leu Gly Gly Asn Val Gly Arg Ile 420 425 430

Ile Leu Glu Asn Asn His Val Ile Asp Thr Tyr Ser Arg Ile Val Phe 435 440 445

Pro Val Val Tyr Ile Ile Phe Asn Leu Phe Tyr Trp Gly Ile Tyr Val 450 455 460

<210> 103

<211> 470

<212> PRT

<213> Morone americana

<400> 103

Met Arg Val Val Leu Leu Ala Leu Arg Leu Met Cys Leu Ala Trp Leu 1 5 10 15

Trp Pro Val Thr Gln Leu Asn Ser Ser Thr Asn Lys Arg Arg His Lys 20 25 30

Glu Leu Tyr Ile Gly Glu Asn Thr Lys Gln Lys His Gly Gly Arg Val $35 \hspace{1.5cm} 40 \hspace{1.5cm} 45$

Asp Leu Lys Leu Lys Lys Val Asp Ser Thr Lys Ser Met Leu Ile Lys 50 60

Ser Glu Gln Leu Leu Arg Ile Glu Asp His Asp Phe Ala Met Arg Pro 65 70 75 80

Gly Phe Gly Gly Ser Ala Ile Pro Val Gly Ile Asp Val Gln Val Glu 85 90 95

Ser Ile Asp Ser Ile Ser Glu Val Asn Met Asp Phe Thr Met Thr Leu Tyr Leu Arg His Tyr Trp Gln Asp Asp Arg Pro Ala Phe Pro Ser Ser Ser Asn Lys Ser Arg Thr Phe Asp Ala Arg Leu Val Lys Ile Trp Val Pro Asp Val Phe Phe Val His Ser Lys Arg Ser Phe Ile His Asp Thr Thr Met Glu Asn Ile Met Leu Arg Val Tyr Pro Asp Gly Asn Ile Leu Tyr Ser Val Arg Ile Thr Val Thr Ala Leu Cys Ser Met Asp Phe Ser Ser Phe Pro Leu Asp Thr Gln Asn Cys Ser Leu Glu Leu Glu Ser Tyr Ala Tyr Ala Tyr Asn Glu Asn Asp Leu Cys Ser Thr Gly Arg Thr Gly Thr Ile Pro Leu Arg Thr Asp Glu Ile Val Leu Ser Gln Phe Phe Val Glu Asp Phe Gln Pro Ser Phe Gly Leu Ala Phe Tyr Ser Ser Thr Gly Trp Tyr Asn Arg Leu Tyr Ile Asn Phe Ile Leu Arg Arg His Ile Phe Phe Phe Met Leu Gln Thr Tyr Phe Pro Thr Met Leu Met Val Met Leu Ser Trp Val Ser Phe Trp Ile Asp Arg Ala Val Pro Ala Arg Val Ser Leu Gly Ile Thr Thr Val Leu Thr Met Ser Thr Ile Ile Thr Gly Val Ser Ala Ser Met Pro Gln Val Ser Tyr Val Lys Ala Val Asp Ile Tyr Leu Trp Ala Ser Phe Leu Phe Val Phe Leu Ser Val Ile Glu Tyr

Ala Ala Val Asn Tyr Phe Thr Thr Val Glu Glu Met Lys Lys Leu Lys 355 360 365

Ser Ala Lys Ile Pro Asn Tyr Asn Ala Ser Gln Ala Met Ala Phe Asp 370 375 380

Gly Cys Phe His Asp Asn Glu Ile Asp Leu Thr Ser Phe Pro Glu Val 385 390 395 400

Ser Ser Thr Pro Asn Thr Glu Arg Asn Thr Gln Ser Arg Asn Ser Asn 405 410 415

Ala Ser Ala Pro Thr Glu Gly Thr Arg Leu Arg Arg Lys His Pro Leu 420 425 430

Arg Gln Asn Leu Ser Phe Ile Met Ser Asn Ser Tyr Met Ile Asp Ser 435 440 445

Tyr Ser Arg Val Ile Phe Pro Leu Ala Tyr Leu Leu Phe Asn Ile Ile 450 455 460

Tyr Trp Ser Met Tyr Ala 465 470

<210> 104

<211> 473

<212> PRT

<213> Homo sapiens

<400> 104

Met Arg Phe Gly Ile Phe Leu Leu Trp Trp Gly Trp Val Leu Ala Thr 1 5 10 15

Glu Ser Arg Met His Trp Pro Gly Arg Glu Val His Glu Met Ser Lys
20 25 30

Lys Gly Arg Pro Gln Arg Gln Arg Glu Val His Glu Asp Ala His 35 40 45

Lys Gln Val Ser Pro Ile Leu Arg Arg Ser Pro Asp Ile Thr Lys Ser 50 55 60

Pro Leu Thr Lys Ser Glu Gln Leu Leu Arg Ile Asp Asp His Asp Phe 65 70 75 80

Ser Met Arg Pro Gly Phe Gly Gly Pro Ala Ile Pro Val Gly Val Asp 85 90 95

Val	Gln	Val	Glu 100	Ser	Leu	Asp	Ser	Ile 105	Ser	Glu	Val	Asp	Met 110	Asp	Phe
Thr	Met	Thr 115	Leu	Tyr	Leu	Arg	His 120	Tyr	Trp	Lys	Asp	Glu 125	Arg	Leu	Ser
Phe	Pro 130	Ser	Thr	Asn	Asn	Leu 135	Ser	Met	Thr	Phe	Asp 140	Gly	Arg	Leu	Val
Lys 145	Lys	Ile	Trp	Val	Pro 150	Asp	Met	Phe	Phe	Val 155	His	Ser	Lys	Arg	Ser 160
Phe	Ile	His	Asp	Thr 165	Thr	Thr	Asp	Asn	Val 170	Met	Leu	Arg	Val	Gln 175	Pro
Asp	Gly	Lys	Val 180	Leu	Tyr	Ser	Leu	Arg 185	Val	Thr	Val.	Thr	Ala 190	Met	Cys
Asn	Met	Asp 195	Phe	Ser	Arg	Phe	Pro 200	Leu	Asp	Thr	Gln	Thr 205	Cys	Ser	Leu
Glu	Ile 210	Glu	Ser	Tyr	Ala	Tyr 215	Thr	Glu	Asp	Asp	Leu 220	Met	Leu	Tyr	Trp
Lys 225	Lys	Gly	Asn	Asp	Ser 230	Leu	Lys	Thr	Asp	Glu 235	Arg	Ile	Ser	Leu	Ser 240
Gln	Phe	Leu	Ile	Gln 245	Glu	Phe	His	Thr	Thr 250	Thr	Lys	Leu	Ala	Phe 255	Tyr
Ser	Ser	Thr	Gly 260	Trp	Tyr	Asn	Arg	Leu 265	Tyr	Ile	Asn	Phe	Thr 270	Leu	Arg
Arg	His	Ile 275	Phe	Phe	Phe	Leu	Leu 280	Gln	Thr	Tyr	Phe	Pro 285	Ala	Thr	Leu
Met	Val 290	Met	Leu	Ser	Trp	Val 295	Ser	Phe	Trp	Ile	Asp 300	Arg	Arg	Ala	Val
Pro 305	Ala	Arg	Val	Pro	Leu 310	Gly	Ile	Thr	Thr	Val 315	Leu	Thr	Met	Ser	Thr 320
Ile	Ile	Thr	Gly	Val 325	Asn	Ala	Ser	Met	Pro 330	Arg	Val	Ser	Tyr	Ile 335	Lys
Ala	Val	Asp	Ile 340	Tyr	Leu	Trp	Val	Ser 345	Phe	Val	Phe	Val	Phe 350	Leu	Ser

Val Leu Glu Tyr Ala Ala Val Asn Tyr Leu Thr Thr Val Gln Glu Arg 355 360 365

Lys Glu Gln Lys Leu Arg Glu Lys Leu Pro Cys Thr Ser Gly Leu Pro 370 375 380

Pro Pro Arg Thr Ala Met Leu Asp Gly Asn Tyr Ser Asp Gly Glu Val 385 390 395 400

Asn Asp Leu Asp Asn Tyr Met Pro Glu Asn Gly Glu Lys Pro Asp Arg
405 410 415

Met Met Val Gln Leu Thr Leu Ala Ser Glu Arg Ser Ser Pro Gln Arg 420 425 430

Lys Ser Gln Arg Ser Ser Tyr Val Ser Met Arg Ile Asp Thr His Ala 435 440 445

Ile Asp Lys Tyr Ser Arg Ile Ile Phe Pro Ala Ala Tyr Ile Leu Phe 450 455 460

Asn Leu Ile Tyr Trp Ser Ile Phe Ser 465 470

<210> 105

<211> 474

<212> PRT

<213> Rattus norvegicus

<400> 105

Met Lys Phe Gly Ile Phe Leu Leu Trp Trp Gly Trp Val Leu Ala Ala 1 5 10 15

Glu Ser Thr Val His Trp Pro Gly Arg Glu Val His Glu Pro Ser Lys
20 25 30

Lys Gly Ser Arg Pro Gln Arg Gln Arg Gly Ala His Asp Asp Ala 35 40 45

His Lys Gln Gly Ser Pro Ile Leu Lys Arg Ser Ser Asp Ile Thr Lys
50 55 60

Ser Pro Leu Thr Lys Ser Glu Gln Leu Leu Arg Ile Asp Asp His Asp 65 70 75 80

Phe Ser Met Arg Pro Gly Phe Gly Gly Pro Ala Ile Pro Val Gly Val

Asp	Val	Gln	Val 100	Glu	Ser	Leu	Asp	Ser 105	Ile	Ser	Glu	Val	Asp 110	Met	Asp
Phe	Thr	Met 115	Thr	Leu	Tyr	Leu	Arg 120	His	Tyr	Trp	Lys	Asp 125	Glu	Arg	Leu
Ser	Phe 130	Pro	Ser	Thr	Asn	Asn 135	Leu	Ser	Met	Thr	Phe 140	Asp	Gly	Arg	Leu
Val 145	Lys	Lys	Ile	Trp	Val 150	Pro	Asp	Met	Phe	Phe 155	Val	His	Ser	Lys	Arg 160
Ser	Phe	Ile	His	Asp 165	Thr	Thr	Thr	Asp	Asn 170	Val	Met	Leu	Arg	Val 175	Gln
Pro	Asp	Gly	Lys 180	Val	Leu	Tyr	Ser	Leu 185	Arg	Val	Thr	Val	Thr 190	Ala	Met
Cys	Asn	Met 195	Asp	Phe	Ser	Arg	Phe 200	Pro	Leu	Asp	Thr	Gln 205	Thr	Cys	Ser
Leu	Glu 210	Ile	Glu	Ser	Tyr	Ala 215	Tyr	Thr	Glu	Asp	Asp 220	Leu	Met	Leu	Tyr
Trp 225	Lys	Lys	Gly	Asn	Asp 230	Ser	Leu	Lys	Thr	Asp 235	Glu	Arg	Ile	Ser	Leu 240
Ser	Gln	Phe	Leu	Ile 245	Gln	Glu	Phe	His	Thr 250	Thr	Thr	Lys	Leu	Ala 255	Phe
Tyr	Ser	Ser	Thr 260	Gly	Trp	Tyr	Asn	Arg 265	Leu	Tyr	Ile	Asn	Phe 270	Thr	Leu
Arg	Arg	His 275	Ile	Phe	Phe	Phe	Leu 280	Leu	Gln	Thr	Tyr	Phe 285	Pro	Ala	Thr
Leu	Met 290	Val	Met	Leu	Ser	Trp 295	Val	Ser	Phe	Trp	Ile 300	Asp	Arg	Arg	Ala
Val 305	Pro	Ala	Arg	Val	Pro 310	Leu	Gly	Ile	Thr	Thr 315	Val	Leu	Thr	Met	Ser 320
Thr	Ile	Ile	Thr	Gly 325	Val	Asn	Ala	Ser	Met 330	Pro	Arg	Val	Ser	Tyr 335	Ile

Lys Ala Val Asp Ile Tyr Leu Trp Val Ser Phe Val Phe Val Phe Leu

340 345 350

Ser Val Leu Glu Tyr Ala Ala Val Asn Tyr Leu Thr Thr Val Gln Glu 355 360 365

Arg Lys Glu Arg Lys Leu Arg Glu Lys Ile Ser Cys Thr Cys Gly Leu 370 380

Pro Gln Pro Arg Gly Val Met Leu Asp Ser Ser Tyr Ser Asp Gly Glu 385 390 395 400

Val Asn Asp Leu Gly Gly Tyr Met Pro Glu Asn Gly Glu Lys Pro Asp 405 410 415

Arg Met Met Val Gln Leu Thr Leu Ala Ser Glu Arg Gly Ser Pro Gln 420 425 430

Arg Lys Ser Gln Arg Gly Ser Tyr Val Ser Met Arg Ile Asn Thr His 435 440 445

Ala Ile Asp Lys Tyr Ser Arg Ile Ile Phe Pro Ala Ala Tyr Ile Leu 450 455 460

Phe Asn Leu Ile Tyr Trp Ser Ile Phe Ser 465 470

<210> 106

<211> 474

<212> PRT

<213> Mus musculus

<400> 106

Met Lys Phe Gly Ile Phe Leu Leu Trp Trp Gly Trp Val Leu Ala Ala 1 5 10 15

Glu Ser Thr Ala His Trp Pro Gly Arg Glu Val His Glu Pro Ser Arg
20 25 30

Lys Gly Ser Arg Pro Gln Arg Gln Arg Gly Ala His Asp Asp Ala 35 40 45

His Lys Gln Gly Ser Pro Ile Leu Arg Arg Ser Ser Asp Ile Thr Lys
50 55 60

Ser Pro Leu Thr Lys Ser Glu Gln Leu Leu Arg Ile Asp Asp His Asp 65 70 75 80

Phe S	Ser	Met	Arg	Pro 85	Gly	Phe	Gly	Gly	Pro 90	Ala	Ile	Pro	Val	Gly 95	Val
Asp '	Val	Gln	Val 100	Glu	Ser	Leu	Asp	Ser 105	Ile	Ser	Glu	Val	Asp 110	Met	Asp
Phe !	Thr	Met 115	Thr	Leu	Tyr	Leu	Arg 120	His	Tyr	Trp	Lys	Asp 125	Glu	Arg	Leu
Ser 1	Phe 130	Pro	Ser	Ser	Asn	Asn 135	Leu	Ser	Met	Thr	Phe 140	Asp	Gly	Arg	Leu
Val 1 145	Lys	Lys	Ile	Trp	Val 150	Pro	Asp	Met	Phe	Phe 155	Val	His	Ser	Lys	Arg 160
Ser 1	Phe	Ile	His	Asp 165	Thr	Thr	Thr	Asp	Asn 170	Val	Met	Leu	Arg	Val 175	Gln
Pro A	Asp	Gly	Lys 180	Val	Leu	Tyr	Ser	Leu 185	Arg	Val	Thr	Val	Thr 190	Ala	Met
Cys I	Asn	Met 195	Asp	Phe	Ser	Arg	Phe 200	Pro	Leu	Asp	Thr	Gln 205	Thr	Суз	Ser
Leu (Glu 210	Ile	Glu	Ser	Tyr	Ala 215	Tyr	Thr	Glu	Asp	Asp 220	Leu	Met	Leu	Tyr
Trp 1 225	Lys	Lys	Gly	Asn	Asp 230	Ser	Leu	Lys	Thr	Asp 235	Glu	Arg	Ile	Ser	Leu 240
Ser (Gln	Phe	Leu	Ile 245	Gln	Glu	Phe	His	Thr 250	Thr	Thr	Lys	Leu	Ala 255	Phe
Tyr S	Ser	Ser	Thr 260	Gly	Trp	Tyr	Asn	Arg 265	Leu	Tyr	Ile	Asn	Phe 270	Thr	Leu
Arg A	Arg	His 275	Ile	Phe	Phe	Phe	Leu 280	Leu	Gln	Thr	Tyr	Phe 285	Pro	Ala	Thr
Leu 1	Met 290	Val	Met	Leu	Ser	Trp 295	Val	Ser	Phe	Trp	Ile 300	Asp	Arg	Arg	Ala
Val 1 305	Pro	Ala	Arg	Val	Pro 310	Leu	Gly	Ile	Thr	Thr 315	Val	Leu	Thr	Met	Ser 320
Thr :	Ile	Ile	Thr	Gly 325	Val	Asn	Ala	Ser	Met 330	Pro	Arg	Val	Ser	Tyr 335	Ile

Lys Ala Val Asp Ile Tyr Leu Trp Val Ser Phe Val Phe Val Phe Leu 340 345 350

Ser Val Leu Glu Tyr Ala Ala Val Asn Tyr Leu Thr Thr Val Gln Glu 355 360 365

Arg Lys Glu Arg Lys Leu Arg Glu Lys Ile Ser Cys Thr Cys Gly Leu 370 375 380

Pro Gln Pro Arg Gly Val Met Leu Asp Ser Ser Tyr Ser Asp Gly Glu 385 390 395 400

Val Asn Asp Leu Gly Gly Tyr Leu Pro Glu Asn Gly Glu Lys Pro Asp 405 410 415

Arg Met Met Val Gln Leu Thr Leu Ala Ser Glu Arg Gly Ser Pro Gln
420 425 430

Arg Lys Gly Gln Arg Gly Ser Tyr Val Ser Met Arg Ile Asn Thr His 435 440 445

Ala Ile Asp Lys Tyr Ser Arg Ile Ile Phe Pro Ala Ala Tyr Ile Leu 450 455 460

Phe Asn Leu Ile Tyr Trp Ser Ile Phe Ser 465 470

<210> 107

<211> 86

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Neur_Chan_LBD domain sequence

<400> 107

Asp Lys Arg Val Arg Pro Val Asn Gly Gly Asp Val Pro Pro Val Thr 1 5 10 15

Val Ser Val Gly Leu Thr Leu Gln Gln Ile Ile Ser Val Asp Glu Lys 20 25 30

Asn Gln Asp Leu Thr Thr Asn Val Trp Leu Arg Gln Gly Gln Trp Thr 35 40 45

Asp Pro Arg Leu Ala Trp Asn Pro Ser Asp Pro Leu Asp Asp Glu Gly

50

55

60

Asp Tyr Gly Gly Ile Lys Ser Leu Arg Leu Pro Ser Asp Asp Asn His 65 70 75 80

Asp Met Leu Asp Lys Ile 85

<210> 108

<211> 67

<212> PRT

<213> Homo sapiens

<400> 108

Asp Phe Ala Met Arg Pro Gly Phe Gly Gly Ser Pro Val Pro Val Gly
1 5 10 15

Ile Asp Val His Val Glu Ser Ile Asp Ser Ile Ser Glu Thr Asn Met 20 25 30

Asp Phe Thr Met Thr Phe Tyr Leu Arg His Tyr Trp Lys Asp Glu Arg 35 40 45

Leu Ser Phe Pro Ser Thr Ala Asn Lys Ser Met Thr Phe Asp His Arg 50 55 60

Lys Ser Ile 65

<210> 109

<211> 25

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
 Neurotransmitter-gated ion-channel domain consensus pattern

<400> 109

Cys Xaa Leu Ile Val Met Phe Gln Xaa Leu Ile Val Met Phe Xaa Xaa 1 5 10 15

Phe Tyr Pro Xaa Asp Xaa Xaa Xaa Cys 20 25 <210> 110

<211> 1015

<212> PRT

<213> Homo sapiens

<400> 110

Met Arg Arg Phe Leu Arg Pro Gly His Asp Pro Val Arg Glu Arg Leu

1 5 10 15

Lys Arg Asp Leu Phe Gln Phe Asn Lys Thr Val Glu His Gly Phe Pro 20 25 30

His Gln Pro Ser Ala Leu Gly Tyr Ser Pro Ser Leu His Ile Leu Ala 35 40 45

Ile Gly Thr Arg Ser Gly Ala Ile Lys Leu Tyr Gly Ala Pro Gly Val 50 55 60

Glu Phe Met Gly Leu His Gln Glu Asn Asn Ala Val Thr Gln Ile His
65 70 75 80

Leu Leu Pro Gly Gln Cys Gln Leu Val Thr Leu Leu Asp Asp Asn Ser 85 90 95

Leu His Leu Trp Ser Leu Lys Val Lys Gly Gly Ala Ser Glu Leu Gln
100 105 110

Glu Asp Glu Ser Phe Thr Leu Arg Gly Pro Pro Gly Ala Ala Pro Ser 115 120 125

Ala Thr Gln Ile Thr Val Val Leu Pro His Ser Ser Cys Glu Leu Leu 130 135 140

Tyr Leu Gly Thr Glu Ser Gly Asn Val Phe Val Val Gln Leu Pro Ala 145 150 155 160

Phe Arg Ala Leu Glu Asp Arg Thr Ile Ser Ser Asp Ala Val Leu Gln 165 170 175

Arg Leu Pro Glu Glu Ala Arg His Arg Arg Val Phe Glu Met Val Glu 180 185 190

Ala Leu Gln Glu His Pro Arg Asp Pro Asn Gln Ile Leu Ile Gly Tyr 195 200 205

Ser Arg Gly Leu Val Val Ile Trp Asp Leu Gln Gly Ser Arg Val Leu 210 215 220

Tyr Hi 225	s Phe	Leu	Ser	Ser 230	Gln	Gln	Leu	Glu	Asn 235	Ile	Trp	Trp	Gln	Arg 240
Asp Gl	y Arg	Leu	Leu 245	Val	Ser	Cys	His	Ser 250	Asp	Gly	Ser	Tyr	Cys 255	Gln
Trp Pr	o Val	Ser 260	Ser	Glu	Ala	Gln	Gln 265	Pro	Glu	Pro	Leu	Arg 270	Ser	Leu
Val Pr	o Tyr 275	Gly	Pro	Phe	Pro	Cys 280	Lys	Ala	Ile	Thr	Arg 285	Ile	Leu	Trp
Leu Th 29		Arg	Gln	Gly	Leu 295	Pro	Phe	Thr	Ile	Phe 300	Gln	Gly	Gly	Met
Pro Ar 305	g Ala	Ser	Tyr	Gly 310	Asp	Arg	His	Cys	Ile 315	Ser	Val	Ile	His	Asp 320
Gly Gl	n Gln	Thr	Ala 325	Phe	Asp	Phe	Thr	Ser 330	Arg	Val	Ile	Gly	Phe 335	Thr
Val Le	u Thr	Glu 340	Ala	Asp	Pro	Ala	Ala 345	Thr	Phe	Asp	Asp	Pro 350	Tyr	Ala
Leu Va	l Val 355	Leu	Ala	Glu	Glu	Glu 360	Leu	Val	Val	Ile	Asp 365	Leu	Gln	Thr
Ala Gl 37		Pro	Pro	Val	Gln 375	Leu	Pro	Tyr	Leu	Ala 380	Ser	Leu	His	Cys
Ser Al 385	a Ile	Thr	Cys	Ser 390	His	His	Val	Ser	Asn 395	Ile	Pro	Leu	Lys	Leu 400
Trp Gl	u Arg	Ile	Ile 405	Ala	Ala	Gly	Ser	Arg 410	Gln	Asn	Ala	His	Phe 415	Ser
Thr Me	t Glu	Trp 420	Pro	Ile	Asp	Gly	Gly 425	Thr	Ser	Leu	Thr	Pro 430	Ala	Pro
Pro Gl	n Arg 435	Asp	Leu	Leu	Leu	Thr 440	Gly	His	Glu	Asp	Gly 445	Thr	Val	Arg
Phe Tr	e Asp	Ala	Ser	Gly	Val 455	Cys	Leu	Arg	Leu	Leu 460	Tyr	Lys	Leu	Ser
Thr Va 465	l Arg	Val	Phe	Leu 470	Thr	Asp	Thr	Asp	Pro 475	Asn	Glu	Asn	Phe	Ser 480

Ala	Gln	Gly	Glu	Asp 485	Glu	Trp	Pro	Pro	Leu 490	Arg	Lys	Val	Gly	Ser 495	Phe
Asp	Pro	Tyr	Ser 500	Asp	Asp	Pro	Arg	Leu 505	Gly	Ile	Gln	Lys	Ile 510	Phe	Leu
Cys	Lys	Tyr 515	Ser	Gly	Tyr	Leu	Ala 520	Val	Ala	Gly	Thr	Ala 525	Gly	Gln	Val
Leu	Val 530	Leu	Glu	Leu	Asn	Asp 535	Glu	Ala	Ala	Glu	Gln 540	Ala	Val	Glu	Gln
Val 545	Glu	Ala	Asp	Leu	Leu 550	Gln	Asp	Gln	Glu	Gly 555	Tyr	Arg	Trp	Lys	Gly 560
His	Glu	Arg	Leu	Ala 565	Ala	Arg	Ser	Gly	Pro 570	Val	Arg	Phe	Glu	Pro 575	Gly
Phe	Gln	Pro	Phe 580	Val	Leu	Val	Gln	Cys 585	Gln	Pro	Pro	Ala	Val 590	Val	Thr
Ser	Leu	Ala 595	Leu	His	Ser	Glu	Trp 600	Arg	Leu	Val	Ala	Phe 605	Gly	Thr	Ser
His	Gly 610	Phe	Gly	Leu	Phe	Asp 615	His	Gln	Gln	Arg	Arg 620	Gln	Val	Phe	Val
Lys 625	Cys	Thr	Leu	His	Pro 630	Ser	Asp	Gln	Leu	Ala 635	Leu	Glu	Gly	Pro	Leu 640
Ser	Arg	Val	Lys	Ser 645	Leu	Lys	Lys	Ser	Leu 650	Arg	Gln	Ser	Phe	Arg 655	Arg
Met	Arg	Arg	Ser 660	Arg	Val	Ser	Ser	Arg 665	Lys	Arg	His	Pro	Ala 670	Gly	Pro
Pro	Gly	Glu 675	Ala	Gln	Glu	Gly	Ser 680	Ala	Lys	Ala	Glu	Arg 685	Pro	Gly	Leu
Gln	Asn 690	Met	Glu	Leu	Ala	Pro 695	Val	Gln	Arg	Lys	Ile 700	Glu	Ala	Arg	Ser
Ala 705	Glu	Asp	Ser	Phe	Thr 710	Gly	Phe	Val	Arg	Thr 715	Leu	Tyr	Phe	Ala	Asp 720
Thr	Tyr	Leu	Lys	Asp 725	Ser	Ser	Arg	His	Cys 730	Pro	Ser	Leu	Trp	Ala 735	Gly

Thr Asn Gly Gly Thr Ile Tyr Ala Phe Ser Leu Arg Val Pro Pro Ala Glu Arg Arg Met Asp Glu Pro Val Arg Ala Glu Gln Ala Lys Glu Ile Gln Leu Met His Arg Ala Pro Val Val Gly Ile Leu Val Leu Asp Gly His Ser Val Pro Leu Pro Glu Pro Leu Glu Val Ala His Asp Leu Ser Lys Ser Pro Asp Met Gln Gly Ser His Gln Leu Leu Val Val Ser Glu Glu Gln Phe Lys Val Phe Thr Leu Pro Lys Val Ser Ala Lys Leu Lys Leu Lys Leu Thr Ala Leu Glu Gly Ser Arg Val Arg Arg Val Ser Val Ala His Phe Gly Ser Arg Arg Ala Glu Asp Tyr Gly Glu His His Leu Ala Val Leu Thr Asn Leu Gly Asp Ile Gln Val Val Ser Leu Pro Leu Leu Lys Pro Gln Val Arg Tyr Ser Cys Ile Arg Arg Glu Asp Val Ser Gly Ile Ala Ser Cys Val Phe Thr Lys Tyr Gly Gln Gly Phe Tyr Leu Ile Ser Pro Ser Glu Phe Glu Arg Phe Ser Leu Ser Thr Lys Trp Leu Val Glu Pro Arg Cys Leu Val Asp Ser Ala Glu Thr Lys Asn His Arg Pro Gly Asn Gly Ala Gly Pro Lys Lys Ala Pro Ser Arg Ala Arg Asn Ser Gly Thr Gln Ser Asp Gly Glu Glu Lys Gln Pro Gly Leu Val Met Glu Arg Ala Leu Leu Ser Asp Glu Arg Ala Ala Thr Gly Val His Ile

Glu Pro Pro Trp Gly Ala Ala Ser Ala Met Ala Glu Gln Ser Glu Trp 995 1000 1005

Leu Ser Val Gln Ala Ala Arg 1010 1015

<210> 111

<211> 1027

<212> PRT

<213> Mus musculus

<400> 111

Met Arg Arg Phe Leu Arg Thr Gly His Asp Pro Ala Arg Glu Arg Leu
1 5 10 15

Lys Arg Asp Leu Phe Gln Phe Asn Lys Thr Val Glu His Gly Phe Pro 20 25 30

His Gln Pro Ser Ala Leu Gly Tyr Ser Pro Ser Leu Arg Ile Leu Ala 35 40 45

Ile Gly Thr Arg Ser Gly Ala Val Lys Leu Tyr Gly Ala Pro Gly Val 50 55 60

Glu Phe Met Gly Leu His Lys Glu Asn Asn Ala Val Leu Gln Ile His
65 70 75 80

Phe Leu Pro Gly Gln Cys Gln Leu Val Thr Leu Leu Asp Asp Asn Ser 85 90 95

Leu His Leu Trp Ser Leu Lys Val Lys Gly Gly Val Ser Glu Leu Gln
100 105 110

Glu Glu Glu Ser Phe Thr Leu Arg Gly Pro Pro Gly Ala Ala Pro Ser 115 120 125

Ala Thr Gln Val Thr Glu Ile Leu Pro His Ser Ser Gly Glu Leu Leu 130 135 140

Tyr Leu Gly Thr Glu Ser Gly Asn Val Leu Val Val Gln Leu Pro Gly
145 150 155 160

Phe Arg Thr Leu His Asp Arg Thr Ile Cys Ser Asp Glu Val Leu Gln
165 170 175

Trp Leu Pro Glu Glu Ala Arg His Arg Arg Val Phe Glu Met Val Glu

Ala	Leu	Gln 195	Glu	His	Pro	Arg	Asp 200	Pro	Asn	Gln	Ile	Leu 205	Ile	Gly	Tyr
Ser	Arg 210	Gly	Leu	Val	Val	Ile 215	Trp	Asp	Leu	Gln	Gly 220	Ser	Arg	Ala	Leu
Ser 225	His	Phe	Leu	Ser	Ser 230	Gln	Gln	Leu	Glu	Asn 235	Ala	Ser	Trp	Gln	Arg 240
Asp	Gly	Cys	Leu	Ile 245	Val	Thr	Cys	His	Ser 250	Asp	Gly	Ser	His	Cys 255	Gln
Trp	Pro	Val	Ser 260	Ser	Asp	Thr	Gln	Asn 265	Pro	Glu	Pro	Leu	Arg 270	Ser	Ser
Ile	Pro	Tyr 275	Gly	Pro	Phe	Pro	Cys 280	Lys	Ala	Ile	Thr	Lys 285	Ile	Phe	Trp
Leu	Thr 290	Thr	Arg	Gln	Gly	Leu 295	Pro	Phe	Thr	Ile	Phe 300	Gln	Gly	Gly	Met
Pro 305	Arg	Ala	Ser	Tyr	Gly 310	Asp	Arg	Asn	Cys	Ile 315	Ser	Val	Val	His	Asn 320
Gly	Gln	Gln	Thr	Gly 325	Phe	Asp	Phe	Thr	Ser 330	Arg	Val	Ile	Asp	Phe 335	Thr
Val	Leu	Ser	Glu 340	Ala	Asp	Pro	Ala	Ala 345	Ala	Phe	Asp	Asp	Pro 350	Tyr	Ala
Leu	Val	Val 355	Leu	Ala	Glu	Glu	Glu 360	Leu	Val	Val	Ile	Asp 365	Leu	Gln	Thr
Pro	Gly 370	Trp	Pro	Pro	Val	Gln 375	Leu	Pro	Tyr	Leu	Ala 380	Ser	Leu	His	Cys
Ser 385	Ala	Ile	Thr	Cys	Ser 390	His	His	Val	Ser	Asn 395	Ile	Pro	Leu	Lys	Leu 400
Trp	Glu	Arg	Ile	Ile 405	Ala	Ala	Gly	Ser	Arg 410	Gln	Asn	Ser	His	Phe 415	Ser
Thr	Met	Glu	Trp 420	Pro	Ile	Asp	Gly	Gly 425	Thr	Ser	Leu	Ala	Pro 430	Pro	Pro

Pro Gln Arg Asp Leu Leu Thr Gly His Glu Asp Gly Thr Val Arg

Phe	Trp 450	Asp	Ala	Ser	Gly	Val 455	Cys	Leu	Arg	Leu	Leu 460	Tyr	Lys	Leu	Ser
Thr 465	Val	Arg	Val	Phe	Leu 470	Thr	Asp	Thr	Asp	Pro 475	Ser	Glu	Asn	Leu	Ser 480
Ala	Gln	Gly	Glu	Asp 485	Glu	Trp	Pro	Pro	Leu 490	Arg	Lys	Val	Gly	Ser 495	Phe
Asp	Pro	Tyr	Ser 500	Asp	Asp	Pro	Arg	Leu 505	Gly	Ile	Gln	Lys	Ile 510	Phe	Leu
Cys	Lys	Tyr 515	Ser	Gly	Tyr	Leu	Ala 520	Val	Ala	Gly	Thr	Ala 525	Gly	Gln	Val
Leu	Val 530	Leu	Glu	Leu	Asn	Asp 535	Glu	Ala	Ala	Glu	His 540	Ala	Val	Glu	Gln
Val 545	Glu	Ala	Asp	Leu	Leu 550	Gln	Asp	Gln	Glu	Gly 555	Tyr	Arg	Trp	Lys	Gly 560
His	Glu	Arg	Leu	Ala 565	Ala	Arg	Pro	Gly	Pro 570	Val	Cys	Phe	Glu	Ala 575	Gly
Phe	Gln	Pro	Phe 580	Val	Leu	Val	Gln	Cys 585	Gln	Pro	Pro	Ala	Val 590	Val	Thr
Ser	Leu	Ala 595	Leu	His	Ser	Glu	Trp 600	Arg	Leu	Val	Ala	Phe 605	Gly	Thr	Ser
His	Gly 610	Phe	Gly	Leu	Phe	Asp 615	His	Gln	Gln	Arg	Arg 620	Gln	Val	Phe	Val
Lys 625	Cys	Thr	Leu	His	Pro 630	Ser	Asp	Gln	Leu	Ala 635	Leu	Glu	Gly	Pro	Leu 640
Ser	Arg	Val	Lys	Ser 645	Leu	Lys	Lys	Ser	Leu 650	Arg	Gln	Ser	Phe	Arg 655	Arg
Met	Arg	Arg	Ser 660	Arg	Val	Ser	Ser	His 665	Lys	Arg	Arg	Pro	Gly 670	Gly	Pro
Thr	Gly	Glu 675	Ala	Gln	Ala	Gln	Ala 680	Val	Asn	Thr	Lys	Thr 685	Glu	Arg	Thr
01 -	T	G1	70	N.TL	a 1.	T	70 T	-		~ 1		-	- 1	~1	70.7 _

Gly Leu Gln Asn Met Glu Leu Ala Pro Val Gln Arg Lys Ile Glu Ala

690	695	700
690	695	700

7\ ** ~*	C	77.7	C1	7. ~ . ~	C	Dh o	Пhъ	C1	Dha	77-1	Vaa	⊞h x	T 011	П.т.	٧aa
705	Ser	AId	GLU	Asp	710	File	TIIT	GIY	rne	715	Add	TIIT	ьеu	ıÀī	720
Ala	Asp	Thr	Tyr	Leu 725	Arg	Asp	Ser	Ser	Arg 730	His	Cys	Pro	Ser	Leu 735	Trp
Ala	Gly	Thr	Asn 740	Gly	Ser	Thr	Val	Tyr 745	Ala	Phe	Ser	Leu	Arg 750	Val	Pro
Pro	Ala	Glu 755	Lys	Lys	Ile	Asn	Lys 760	Pro	Val	Arg	Ala	Lys 765	Gln	Ala	Lys
Glu	Ile 770	Gln	Leu	Met	His	Arg 775	Ala	Pro	Val	Val	Gly 780	Ile	Leu	Val	Leu
Asp 785	Gly	His	Asn	Val	Pro 790	Leu	Pro	Glu	Pro	Leu 795	Glu	Val	Ala	His	Asp 800
Leu	Ser	Lys	Ser	Pro 805	Asp	Met	Gln	Gly	Ser 810	His	Gln	Leu	Leu	Val 815	Val
Ser	Glu	Glu	Gln 820	Phe	Lys	Val	Phe	Thr 825	Leu	Pro	Lys	Val	Ser 830	Ala	Lys
Leu	Lys	Leu 835	Lys	Leu	Thr	Ala	Leu 840	Glu	Gly	Ser	Arg	Val 845	Arg	Arg	Val
Gly	Val 850	Ala	His	Phe	Gly	Ser 855	Cys	Arg	Ala	Glu	Asp 860	Tyr	Gly	Glu	His
His 865	Leu	Ala	Val	Leu	Thr 870	Asn	Leu	Gly	Asp	Ile 875	Gln	Val	Val	Ser	Met 880
Pro	Leu	Leu	Lys	Pro 885	Gln	Val	Arg	Tyr		Cys	Ile	Arg	Arg	Glu 895	Asp
Val	Ser	Gly	Ile 900	Ala	Ser	Cys	Val	Phe 905	Thr	Lys	Tyr	Gly	Gln 910	Gly	Phe
Tyr	Leu	Ile 915	Ser	Pro	Ser	Glu	Phe 920	Glu	Arg	Phe	Ser	Leu 925	Ser	Thr	Lys
Trp	Leu 930	Val	Glu	Pro	Arg	Cys 935	Leu	Val	Asp	Ser	Thr	Lys	Ala	Lys	Lys

His Asn Arg Pro Ser Asn Gly Asn Gly Thr Gly Pro Lys Met Thr Ser

945 950 955 960

Ser Gly His Val Arg Asn Ser Lys Ser Gln Ser Asp Gly Asp Glu Lys 965 970 975

Lys Pro Gly Pro Val Met Glu His Ala Leu Leu Asn Asp Ala Trp Val 980 985 990

Leu Lys Glu Ile Gln Ser Thr Leu Glu Gly Asp Arg Arg Ser Tyr Gly 995 1000 1005

Asn Trp His Pro His Arg Val Ala Val Gly Cys Arg Leu Ser Asn Gly 1010 1015 1020

Glu Ala Glu 1025

<210> 112

<211> 1034

<212> PRT

<213> Mus musculus

<400> 112

Met Met Lys Phe Arg Phe Arg Gln Gly Ala Asp Pro Gln Arg Glu 1 5 10 15

Lys Leu Lys Gln Glu Leu Phe Ala Phe His Lys Thr Val Glu His Gly 20 25 30

Phe Pro Asn Gln Pro Ser Ala Leu Ala Phe Asp Pro Glu Leu Arg Ile 35 40 45

Met Ala Ile Gly Thr Arg Ser Gly Ala Val Lys Ile Tyr Gly Ala Pro 50 55 60

Gly Val Glu Phe Thr Gly Leu His Arg Asp Ala Ala Thr Val Thr Gln 65 70 75 80

Met His Phe Leu Pro Gly Gln Gly Arg Leu Leu Thr Leu Leu Asp Asp 85 90 95

Ser Ser Leu His Leu Trp Glu Ile Ile His His Asn Gly Cys Ala His
100 105 110

Leu Glu Glu Gly Leu Ser Phe His Pro Pro Ser Arg Pro Ser Phe Asp 115 120 125

Asn Ala Ser 130	Phe Pro	Ala Ser 135	Leu Thi	r Arg Val	Thr V	al Val	Leu	Leu
Val Ala Gly 145	Asn Thr	Ala Ala 150	Leu Gly	y Thr Glu 155		ly Ser	Ile	Phe 160
Phe Leu Asp	Val Ala 165	Thr Leu	Ala Le	ı Leu Glı 170	ı Gly G	ln Thr	Leu 175	Ser
Pro Asp Val	Val Leu 180	Arg Ser	Val Pro		Tyr A	rg Cys 190	Gly	Lys
Ala Leu Gly 195	Pro Val	Glu Ser	Leu Gli 200	n Gly His		ln Asp 05	Pro	Ser
Lys Ile Leu 210	Ile Gly	Tyr Ser 215	Arg Gly	y Leu Lei	val I 220	le Trp	Ser	Gln
Ala Thr Gln 225		Asp Asn 230	Val Phe	e Leu Gly 235		ln Gln	Leu	Glu 240
Ser Leu Cys	Trp Gly 245	Arg Asp	Gly Se	r Ser Ile 250	e Ile S	er Ser	His 255	Ser
Asp Gly Ser	Tyr Ala 260	Ile Trp	Ser Thi	_	Gly S	er Pro 270	Pro	Thr
Leu Gln Pro 275	Thr Val	Val Thr	Thr Pro	o Tyr Gly		he Pro 85	Cys	Lys
Ala Ile Asn 290	Lys Ile	Leu Trp 295	Arg Sei	c Cys Gli	Ser G 300	ly Asp	His	Phe
Ile Ile Phe 305		Gly Met 310	Pro Arc	g Ala Sei 315		ly Asp	Arg	His 320
Cys Val Ser	Val Leu 325	Arg Ala	Glu Th	r Leu Val 330	Thr L	eu Asp	Phe 335	Thr
Ser Arg Val	Ile Asp 340	Phe Phe	Thr Val		Thr G	ln Pro 350	Glu	Asp
Glu Cys Asp 355	Asn Pro	Gln Ala	Leu Ala	a Val Lei		lu Glu 65	Glu	Leu
Val Val Leu 370	Asp Leu	Gln Thr 375	Pro Gly	y Trp Pro	Ala V 380	al Pro	Ala	Pro

Tyr 385	Leu	Ala	Pro	Leu	His 390	Ser	Ser	Ala	Ile	Thr 395	Cys	Ser	Ala	His	Val 400	
Ala	Asn	Val	Pro	Ser 405	Lys	Leu	Trp	Ala	Arg 410	Ile	Val	Ser	Ala	Gly 415	Glu	
Gln	Gln	Ser	Pro 420	Gln	Pro	Ala	Ser	Ser 425	Ala	Leu	Ser	Trp	Pro 430	Ile	Thr	
Gly	Gly	Arg 435	Asn	Leu	Ala	Gln	Glu 440	Pro	Ser	Gln	Arg	Gly 445	Leu	Leu	Leu	
Thr	Gly 450	His	Glu	Asp	Gly	Thr 455	Val	Arg	Phe	Trp	Asp 460	Ala	Ser	Gly	Val	
Ala 465	Leu	Arg	Pro	Leu	Tyr 470	Lys	Leu	Ser	Thr	Ala 475	Gly	Leu	Phe	Gln	Thr 480	
Asp	Cys	Glu	His	Ala 485	Asp	Ser	Leu	Ala	Gln 490	Ala	Val	Glu	Asp	Asp 495	Trp	
Pro	Pro	Phe	Arg 500	Lys	Val	Gly	Cys	Phe 505	Asp	Pro	Tyr	Ser	Asp 510	Asp	Pro	
Arg	Leu	Gly 515	Ile	Gln	Lys	Val	Ala 520	Leu	Cys	Lys	Tyr	Thr 525	Ala	Gln	Met	
Val	Val 530	Ala	Gly	Thr	Ala	Gly 535	Gln	Val	Leu	Val	Leu 540	Glu	Leu	Ser	Glu	
Val 545	Pro	Ala	Glu	His	Ala 550	Val	Ser	Val	Ala	Asn 555	Val	Asp	Leu	Leu	Gln 560	
Asp	Arg	Glu	Gly	Phe 565	Thr	Trp	Lys	Gly	His 570	Glu	Arg	Leu	Asn	Pro 575	His	
Thr	Gly	Leu	Leu 580	Pro	Trp	Pro	Ala	Gly 585	Phe	Gln	Pro	Arg	Met 590	Leu	Ile	
Gln	Cys	Leu 595	Pro	Pro	Ala	Ala	Val 600	Thr	Ala	Val	Thr	Leu 605	His	Ala	Glu	
Trp	Ser 610	Leu	Val	Ala	Phe	Gly 615	Thr	Ser	His	Gly	Phe 620	Gly	Leu	Phe	Asp	
Tyr 625	Gln	Arg	Lys	Ser	Pro 630	Val	Leu	Ala	Arg	Cys 635	Thr	Leu	His	Pro	Asn 640	

Asp	Ser	Leu	Ala	Met 645	Glu	Gly	Pro	Leu	Ser 650	Arg	Val	Lys	Ser	Leu 655	Lys
Lys	Ser	Leu	Arg 660	Gln	Ser	Phe	Arg	Arg 665	Ile	Arg	Lys	Ser	Arg 670	Val	Ser
Gly	Lys	Lys 675	Arg	Thr	Pro	Ala	Ala 680	Ser	Ser	Lys	Glu	Ala 685	Asn	Ala	Gln
Leu	Ala 690	Glu	Gln	Thr	Cys	Pro 695	His	Asp	Leu	Glu	Met 700	Thr	Pro	Val	Gln
Arg 705	Arg	Ile	Glu	Pro	Arg 710	Ser	Ala	Asp	Asp	Ser 715	Leu	Ser	Gly	Val	Val 720
Arg	Cys	Leu	Tyr	Phe 725	Ala	Asp	Thr	Phe	Leu 730	Arg	Asp	Ala	Thr	His 735	His
Gly	Pro	Thr	Met 740	Trp	Ala	Gly	Thr	Asn 745	Ser	Gly	Ser	Val	Phe 750	Ala	Tyr
Ala	Leu	Glu 755	Val	Pro	Ala	Ala	Thr 760	Ala	Gly	Gly	Glu	Lys 765	Arg	Pro	Glu
Gln	Ala 770	Val	Glu	Ala	Val	Leu 775	Gly	Lys	Glu	Val	Gln 780	Leu	Met	His	Arg
Ala 785	Pro	Val	Val	Ala	Ile 790	Ala	Val	Leu	Asp	Gly 795	Arg	Gly	Arg	Pro	Leu 800
Pro	Glu	Pro	Tyr	Glu 805	Ala	Ser	Arg	Asp	Leu 810	Ala	Gln	Ala	Pro	Asp 815	Met
Gln	Gly	Gly	His 820	Ala	Val	Leu	Ile	Ala 825	Ser	Glu	Glu	Gln	Phe 830	Lys	Val
Phe	Thr	Leu 835	Pro	Lys	Val	Ser	Ala 840	Lys	Thr	Lys	Phe	Lys 845	Leu	Thr	Ala
His	Glu 850	Gly	Cys	Arg	Val	Arg 855	Lys	Val	Ala	Leu	Ala 860	Thr	Phe	Ala	Ser
Val 865	Met	Ser	Glu	Asp	Tyr 870	Ala	Glu	Thr	Cys	Leu 875	Ala	Cys	Leu	Thr	Asn 880
Leu	Gly	Asp	Val	His 885	Val	Phe	Ser	Val	Pro 890	Gly	Leu	Arg	Pro	Gln 895	Val

His Tyr Ser Cys Ile Arg Lys Glu Asp Ile Ser Gly Ile Ala Ser Cys 900 905 910

Val Phe Thr Arg His Gly Gln Gly Phe Tyr Leu Ile Ser Pro Ser Glu 915 920 925

Phe Glu Arg Phe Ser Leu Ser Ala Arg Asn Ile Thr Glu Pro Leu Cys 930 935 940

Ser Leu Asp Ile Ser Trp Pro Gln Asn Ala Thr Gln Pro Arg Leu Gln 945 950 955 960

Glu Ser Pro Lys Leu Ser Gln Ala Asn Gly Thr Arg Asp Ile Ile Leu 965 970 975

Ala Pro Glu Ser Cys Glu Gly Ser Pro Ser Ser Ala His Ser Lys Arg 980 985 990

Ala Asp Thr Met Glu Pro Pro Glu Ala Ala Leu Ser Pro Val Ser Ile 995 1000 1005

Asp Ser Ala Ala Ser Gly Asp Thr Met Leu Asp Thr Thr Gly Asp Val 1010 1015 1020

Thr Val Glu Tyr Val Lys Asp Phe Leu Gly 1025 1030

<210> 113

<211> 1057

<212> PRT

<213> Homo sapiens

<400> 113

Met Met Lys Phe Pro Phe Arg Arg Gln Gly Ala Asp Pro Gln Arg Glu
1 5 10 15

Lys Leu Lys Gln Glu Leu Phe Ala Phe Asn Lys Thr Val Glu His Gly 20 25 30

Phe Pro Asn Gln Pro Ser Ala Leu Ala Phe Asp Pro Glu Leu Arg Ile 35 40 45

Met Ala Ile Gly Thr Arg Ser Gly Ala Val Lys Ile Tyr Gly Ala Pro 50 55 60

Gly Val Glu Phe Thr Gly Leu His Arg Asp Ala Ala Thr Val Thr Gln
65 70 75 80

Met	His	Phe	Leu	Thr 85	Gly	Gln	Gly	Arg	Leu 90	Leu	Ser	Leu	Leu	Asp 95	Asp
Ser	Ser	Leu	His 100	Leu	Trp	Glu	Ile	Val 105	His	His	Asn	Gly	Cys 110	Ala	His
Leu	Glu	Glu 115	Ala	Leu	Ser	Phe	Gln 120	Leu	Pro	Ser	Arg	Pro 125	Gly	Phe	Asp
Gly	Ala 130	Ser	Ala	Pro	Leu	Ser 135	Leu	Thr	Arg	Val	Thr 140	Val	Val	Leu	Leu
Val 145	Ala	Ala	Gly	Asp	Ile 150	Ala	Ala	Leu	Gly	Thr 155	Glu	Gly	Ser	Ser	Ser 160
Val	Phe	Phe	Leu	Asp 165	Val	Thr	Thr	Leu	Thr 170	Leu	Leu	Glu	Gly	Gln 175	Thr
Leu	Ala	Pro	Gly 180	Glu	Val	Leu	Arg	Ser 185	Val	Pro	Asp	Asp	Tyr 190	Arg	Cys
Gly	Lys	Asp 195	Leu	Gly	Pro	Val	Glu 200	Ser	Leu	Gln	Gly	His 205	Leu	Gln	Asp
Pro	Thr 210	Lys	Ile	Leu	Ile	Gly 215	Tyr	Ser	Arg	Gly	Leu 220	Leu	Val	Ile	Arg
Asn 225	Gln	Ala	Ser	Gln	Cys 230	Val	Asp	His	Ile	Phe 235	Leu	Gly	Asn	Gln	Gln 240
Leu	Glu	Ser	Leu	Cys 245	Trp	Gly	Arg	Asp	Ser 250	Ser	Thr	Val	Val	Ser 255	Ser
His	Ser	Asp	Gly 260	Ser	Tyr	Ala	Val	Trp 265	Ser	Val	Asp	Ala	Gly 270	Ser	Phe
Pro	Thr	Leu 275	Gln	Pro	Thr	Val	Ala 280	Thr	Thr	Pro	Tyr	Gly 285	Pro	Phe	Pro
Cys	Lys 290	Ala	Ile	Asn	Lys	Ile 295	Leu	Trp	Arg	Asn	Cys 300	Glu	Ser	Gly	Gly
His 305	Phe	Ile	Ile	Phe	Ser 310	Gly	Gly	Met	Pro	Arg 315	Ala	Ser	Tyr	Gly	Asp 320
Arg	His	Cys	Val	Ser 325	Val	Leu	Arg	Ala	Glu 330	Thr	Leu	Val	Thr	Leu 335	Asp

Phe His Phe Arg Ile Ile Asp Phe Phe Thr Val His Ser Thr Arg Pro Glu Asp Glu Phe Asp Asp Pro Gln Ala Leu Ala Val Leu Leu Glu Glu Glu Leu Val Val Leu Asp Leu Gln Thr Pro Gly Trp Pro Ala Val Pro Ala Pro Tyr Leu Ala Pro Leu His Ser Ser Ala Ile Thr Cys Ser Ala His Val Ala Ser Val Pro Ala Lys Leu Trp Ala Arg Ile Val Ser Ala Gly Glu Gln Ser Pro Gln Pro Val Ser Ser Ala Leu Ser Trp Pro Ile Thr Gly Gly Arg Asn Leu Ala Gln Glu Pro Ser Gln Arg Gly Leu Leu Leu Thr Gly His Glu Asp Gly Thr Val Arg Phe Trp Asp Ala Ser Gly Val Ala Leu Arg Pro Leu Tyr Lys Leu Ser Thr Ala Gly Leu Phe Gln Thr Asp Cys Glu His Ser Asp Ser Leu Ala Gln Ala Ala Glu Asp Asp Trp Pro Pro Phe Arg Lys Val Gly Cys Phe Asp Pro Tyr Ser Asp Asp Pro Arg Leu Gly Val Gln Lys Val Ala Leu Cys Lys Tyr Thr Ala Gln Met Val Val Ala Gly Thr Ala Gly Gln Val Leu Val Leu Glu Leu Ser Asp Val Pro Val Glu His Ala Val Ser Val Ala Ile Ile Asp Leu Leu Gln Asp Arg Glu Gly Phe Thr Trp Lys Gly His Glu Arg Leu Ser Pro Arg Thr Gly Leu Leu Pro Trp Pro Ala Gly Phe Gln Pro Cys Val

Leu Val Gln Cys Leu Pro Pro Ala Ala Val Thr Ala Val Thr Leu His Thr Glu Trp Ser Leu Val Ala Phe Gly Thr Ser His Gly Phe Gly Leu Leu Ser Pro Val Leu Ala Arg Cys Thr Leu His Pro Asn Asp Ser Leu Ala Met Glu Gly Pro Leu Ser Arg Val Lys Ser Leu Lys Lys Ser Leu Arg Gln Ser Phe Arg Arg Ile Arg Lys Ser Arg Val Ser Gly Lys Lys Arg Ala Ala Asn Ala Ser Ser Lys Leu Gln Glu Ala Asn Ala Gln Leu Ala Glu Gln Ala Cys Pro His Asp Val Glu Met Thr Pro Val Gln Arg Arg Ile Glu Pro Arg Ser Ala Asp Asp Ser Leu Ser Gly Val Val Arg Cys Leu Tyr Phe Ala Asp Thr Phe Leu Arg Asp Gly Ala His His Gly Pro Thr Met Trp Ala Gly Thr Asn Ser Gly Ser Val Phe Ala Tyr Ala Leu Glu Val Pro Ala Ala Ala Val Gly Glu Lys Arg Pro Glu Gln Ala Val Glu Ala Val Leu Gly Lys Glu Leu Gln Leu Met His Arg Ala Pro Val Val Ala Ile Ala Val Leu Asp Gly Gly Arg Pro Leu Pro Glu Pro Tyr Glu Ala Ser Arg Asp Leu Ala Gln Ala Pro His Met Gln Gly Gly His Ala Val Leu Ile Ala Ser Glu Glu Gln Phe Lys Val Phe Thr Leu Pro Lys Val Ser Ala Lys Thr Lys Phe Lys Leu Thr Ala His Glu

Gly Cys Arg Val Arg Lys Val Val Ala Leu Ala Thr Phe Ala Ser Val 850 855 860

Ala Cys Glu Asp Tyr Ala Glu Thr Cys Leu Ala Cys Leu Thr Asn Leu 865 870 875 880

Gly Asp Val His Val Phe Ser Val Pro Gly Leu Arg Pro Glu Val His 885 890 895

Tyr Ser Cys Ile Arg Lys Glu Asp Ile Ser Gly Ile Ala Ser Cys Val 900 905 910

Phe Thr Arg His Gly Gln Gly Phe Tyr Leu Ile Ser Pro Ser Glu Phe 915 920 925

Glu Arg Phe Ser Leu Ser Ala Arg Asn Ile Thr Glu Gly Leu Cys Ser 930 935 940

Leu Asp Ile Asn Trp Pro Arg Asp Ala Thr Gln Ala Ser Tyr Arg Ile 945 950 955 960

Arg Glu Ser Pro Lys Leu Ser Gln Ala Asn Gly Thr Pro Ser Ile Leu 965 970 975

Leu Ala Pro Gln Ser Leu Asp Gly Ser Pro Asp Pro Ala His Ser Met 980 985 990

Gly Pro Asp Thr Pro Glu Pro Pro Glu Ala Ala Leu Ser Pro Met Ser 995 1000 1005

Ile Asp Ser Ala Thr Ser Ala Asp Thr Thr Leu Asp Thr Thr Gly Asp 1010 1015 1020

Val Thr Val Glu Asp Val Lys Asp Phe Leu Gly Ser Ser Glu Glu Ser 1025 1030 1035 1040

Glu Lys Asn Leu Arg Asn Leu Ala Glu Asp Glu Ala His Ala Cys Cys 1045 1050 1055

Ile

<210> 114

<211> 1032

<212> PRT

<213> Homo sapiens

- Met Met Lys Phe Arg Phe Arg Gln Gly Ala Asp Pro Gln Arg Glu
 1 5 10 15
- Lys Leu Lys Gln Glu Leu Phe Ala Phe Asn Lys Thr Val Glu His Gly
 20 25 30
- Phe Pro Asn Gln Pro Ser Ala Leu Ala Phe Asp Pro Glu Leu Arg Ile 35 40 45
- Met Ala Ile Gly Thr Arg Ser Gly Ala Val Lys Ile Tyr Gly Ala Pro 50 55 60
- Gly Val Glu Phe Thr Gly Leu His Arg Asp Ala Ala Thr Val Thr Gln 65 70 75 80
- Met His Phe Leu Thr Gly Gln Gly Arg Leu Leu Ser Leu Leu Asp Asp 85 90 95
- Ser Ser Leu His Leu Trp Glu Ile Val His His Asn Gly Cys Ala His
 100 105 110
- Leu Glu Glu Ala Leu Ser Phe Gln Leu Pro Ser Arg Pro Gly Phe Asp 115 120 125
- Gly Ala Ser Ala Pro Leu Ser Leu Thr Arg Val Thr Val Val Leu Leu 130 135 140
- Val Ala Ala Gly Asp Ile Ala Gly Leu Gly Thr Glu Gly Ser Ser Val 145 150 155 160
- Phe Phe Leu Asp Val Thr Thr Leu Thr Leu Glu Gly Gln Thr Leu
 165 170 175
- Ala Pro Gly Glu Val Leu Arg Ser Val Pro Asp Asp Tyr Arg Cys Gly
 180 185 190
- Lys Ala Leu Gly Pro Val Glu Ser Leu Gln Gly His Cys Gly Thr Pro
 195 200 205
- Gln Arg Phe Ser Leu Ala Thr Asp Arg Gly Leu Leu Val Ile Trp Asn 210 215 220
- Gln Ser Arg Gln Cys Val Asp His Ile Phe Leu Gly Asn Gln Gln Leu 225 230 235 240
- Glu Ser Leu Cys Trp Gly Arg Asp Ser Ser Thr Val Val Ser Ser His

				243					230					255	
Ser	Asp	Gly	Ser 260	Tyr	Ala	Val	Trp	Ser 265	Val	Asp	Ala	Gly	Ser 270	Phe	Pro
Thr	Leu	Gln 275	Pro	Thr	Val	Ala	Thr 280	Thr	Pro	Tyr	Gly	Pro 285	Phe	Pro	Cys
Lys	Ala 290	Ile	Asn	Lys	Ile	Leu 295	Trp	Arg	Asn	Cys	Glu 300	Ser	Gly	Gly	His
Phe 305	Ile	Ile	Phe	Ser	Gly 310	Gly	Met	Pro	Arg	Ala 315	Ser	Tyr	Gly	Asp	Arg 320
His	Cys	Val	Ser	Val 325	Leu	Arg	Ala	Glu	Thr 330	Leu	Val	Thr	Leu	Asp 335	Phe
Thr	Ser	Arg	Ile 340	Ile	Asp	Phe	Phe	Thr 345	Val	His	Ser	Thr	Arg 350	Pro	Glu
Asp	Glu	Phe 355	Asp	Asp	Pro	Gln	Ala 360	Leu	Ala	Val	Leu	Leu 365	Glu	Glu	Glu
Leu	Val 370	Val	Leu	Asp	Leu	Gln 375	Thr	Pro	Gly	Trp	Pro 380	Ala	Val	Pro	Ala
Pro 385	Tyr	Leu	Ala	Pro	Leu 390	His	Ser	Ser	Ala	Ile 395	Thr	Cys	Ser	Ala	Tyr 400
Val	Ala	Ser	Val	Pro 405	Ala	Lys	Leu	Trp	Ala 410	Arg	Ile	Val	Ser	Ala 415	Gly
Glu	Gln	Gln	Ser 420	Pro	Gln	Pro	Val	Ser 425	Ser	Ala	Leu	Ser	Trp 430	Pro	Ile
Thr	Gly	Gly 435	Arg	Asn	Leu	Ala	Gln 440	Glu	Pro	Ser	Gln	Arg 445	Gly	Leu	Leu
Leu	Thr 450	Gly	His	Glu	Asp	Gly 455	Thr	Val	Arg	Phe	Trp 460	Asp	Ala	Ser	Gly
Val 465	Ala	Leu	Arg	Pro	Leu 470	Tyr	Lys	Leu	Ser	Thr 475	Ala	Gly	Leu	Phe	Gln 480
Thr	Asp	Cys	Glu	His 485	Ala	Asp	Ser	Leu	Ala 490	Gln	Ala	Ala	Glu	Asp 495	Asp

Trp Pro Pro Phe Arg Lys Val Gly Cys Phe Asp Pro Tyr Ser Asp Asp

Pro	Arg	Leu 515	Gly	Val	Gln	Lys	Val 520	Ala	Leu	Cys	Lys	Tyr 525	Thr	Ala	Gln
Met	Val 530	Val	Ala	Gly	Thr	Ala 535	Gly	Gln	Val	Leu	Val 540	Leu	Glu	Leu	Ser
Asp 545	Val	Pro	Val	Glu	Gln 550	Ala	Val	Ser	Val	Ala 555	Ile	Ile	Asp	Leu	Leu 560
Gln	Asp	Arg	Glu	Gly 565	Phe	Thr	Trp	Lys	Gly 570	His	Glu	Arg	Leu	Ser 575	Pro
Arg	Thr	Gly	Pro 580	Leu	Pro	Trp	Pro	Ala 585	Gly	Phe	Leu	Pro	Arg 590	Val	Leu
Val	Gln	Cys 595	Leu	Pro	Pro	Ala	Ala 600	Val	Thr	Ala	Val	Thr 605	Leu	His	Thr
Glu	Trp 610	Ser	Leu	Val	Ala	Phe 615	Gly	Thr	Ser	His	Gly 620	Phe	Gly	Leu	Phe
Asp 625	Tyr	Gln	Arg	Lys	Ser 630	Pro	Val	Leu	Ala	Arg 635	Cys	Thr	Leu	His	Pro 640
Asn	Asp	Ser	Leu	Ala 645	Met	Glu	Gly	Pro	Leu 650	Ser	Arg	Val	Lys	Ser 655	Leu
Lys	Lys	Ser	Leu 660	Arg	Gln	Ser	Phe	Arg 665	Arg	Ile	Arg	Lys	Ser 670	Arg	Val
Ser	Gly	Lys 675	Lys	Arg	Ala	Ala	Asn 680	Ala	Ser	Ser	Lys	Leu 685	Leu	Glu	Ala
Asn	Ala 690	Gln	Leu	Ala	Glu	Gln 695	Ala	Cys	Pro	His	Asp 700	Val	Glu	Met	Thr
Pro 705	Val	Gln	Arg	Arg	Ile 710	Glu	Pro	Arg	Ser	Ala 715	Asp	Asp	Ser	Leu	Ser 720
Gly	Val	Val	Arg	Cys 725	Leu	Tyr	Phe	Ala	Asp 730	Thr	Phe	Leu	Arg	Asp 735	Gly
Pro	Thr	Thr	Gly 740	Pro	Thr	Met	Trp	Ala 745	Gly	Thr	Asn	Ser	Gly 750	Ser	Val

Phe Ala Tyr Ala Leu Glu Val Pro Ala Ala Ala Val Gly Gly Glu Lys

755 760	765
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Ara	Pro	Glu	Gln	Ala	Val	Glu	Ala	Val	Leu	Gly	Lys	Lys	Glu	Gln	Leu
5	770					7 75				•	780	-			
Met 785	His	Arg	Ala	Pro	Val 790	Val	Ala	Ile	Cys	Arg 795	Val	Gly	Arg	Arg	Gly 800
Arg	Pro	Leu	Pro	Glu 805	Pro	Tyr	Glu	Ala	Ser 810	Arg	Asp	Leu	Ala	Gln 815	Ala
Pro	Asp	Met	Gln 820	Gly	Gly	His	Ala	Val 825	Leu	Ile	Ala	Ser	Glu 830	Glu	Gln
Phe	Lys	Val 835	Phe	Thr	Leu	Pro	Lys 840	Val	Ser	Ala	Lys	Thr 845	Lys	Phe	Lys
Leu	Thr 850	Ala	His	Glu	Gly	Cys 855	Arg	Val	Arg	Lys	Val 860	Ala	Leu	Ala	Thr
Phe 865	Cys	Gln	Cys	Gly	Leu 870	Gln	Thr	Met	Leu	Arg 875	Pro	Ala	Trp	Pro	Val 880
Leu	Thr	Asn	Leu	Gly 885	Asp	Val	His	Val	Phe 890	Ser	Val	Pro	Leu	Arg 895	Pro
Gln	Val	His	Tyr 900	Ser	Cys	Ile	Arg	Lys 905	Glu	Asp	Ile	Ser	Gly 910	Ile	Ala
Ser	Cys	Val 915	Phe	Thr	Arg	His	Gly 920	Gln	Gly	Phe	Tyr	Leu 925	Ile	Ser	Pro
Ser	Glu 930	Phe	Glu	Arg	Phe	Ser 935	Leu	Ser	Ala	Arg	Asn 940	Ile	Thr	Glu	Arg
Ser 945	Ala	Leu	Trp	Thr	Leu 950	Thr	Gly	Pro	Ala	Met 955	Pro	Pro	Arg	Pro	Val 960
Thr	Gly	Ser	Glu	Ser 965	His	Pro	Lys	Leu	Ser 970	Gln	Ala	Asn	Gly	Thr 975	Pro
Ser	Ile	Leu	Leu 980	Ala	Pro	Gln	Ser	Leu 985	Asp	Gly	Ser	Pro	Asp 990	Pro	Ala
His	Ser	Met 995	Gly	Pro	Asp		Pro 1000	Glu	Pro	Pro		Ala 1005	Ala	Leu	Ser

Pro Met Ser Ile Asp Ser Ala Thr Ser Ala Asp Thr Thr Leu Thr Arg

1010 1015 1020

Gln Gly Thr Ser Gln Trp Lys Met 1025 1030

<210> 115

<211> 36

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: WD domain sequence

<400> 115

Leu Leu Arg Thr Leu Gly His Ser Ser Ser Val Thr Ser Leu Ala Phe
1 5 10 15

Asp Pro Asp Gly Gly Leu Leu Ala Thr Gly Ser Ala Asp Gly Thr Val
20 25 30

Arg Ile Trp Asp 35

<210> 116

<211> 37

<212> PRT

<213> Homo sapiens

<400> 116

Asn Lys Thr Val Glu His Gly Phe Pro His Gln Pro Ser Ala Leu Gly 1 5 10 15

Tyr Ser Pro Ser Leu His Ile Leu Ala Ile Gly Thr Arg Ser Gly Ala 20 25 30

Ile Lys Leu Tyr Gly 35

<210> 117

<211> 1130

<212> PRT

<213> Homo sapiens

<400> 117

Gly Val Asn Ala Gln Thr Lys Asn Gly Ala Thr Pro Leu Tyr Leu Ala Cys Gln Glu Gly His Leu Glu Val Thr Gln Tyr Leu Val Gln Glu Cys Gly Ala Asp Pro His Ala Arg Ala His Asp Gly Met Thr Pro Leu His Ala Ala Gln Met Gly His Ser Pro Val Ile Val Trp Leu Val Ser Cys Thr Asp Val Ser Leu Ser Glu Gln Asp Lys Asp Gly Ala Thr Ala Thr His Phe Ala Ala Ser Arg Gly His Ser Lys Val Leu Ser Trp Leu Leu Leu His Gly Gly Glu Ile Ser Ala Asp Leu Trp Gly Gly Thr Ala Leu Tyr Asp Ala Ala Glu Asn Gly Glu Leu Glu Cys Cys Gln Ile Leu Val Val Asn Gly Ala Glu Leu Glu Val Arg Asp Arg Asp Gly Tyr Ala Ala Ala Asp Leu Ser Asp Phe Asn Gly His Ser His Cys Thr His Cys Leu Arg Thr Val Glu Asn Leu Ser Met Glu His Cys Val Leu Ser Arg Asp Pro Ser Val Glu Leu Glu Ala Lys Gln Pro Asp Ser Gly Met Ser Ser Pro Asn Thr Thr Val Ser Val Gln Pro Leu Asn Phe Asp Leu Ser Ser Pro Thr Ser Thr Leu Ser Asn Tyr Asp Ser Cys Ser Ser Ser His Ser Ser Ile Lys Gly Gln His Pro Pro Arg Gly Leu Ser Ser Thr Arg Ala Ala Asp Ile Gln Ser Tyr Met Asp Met Leu Asn Pro Glu Leu Gly

Leu P	ro Tr	260	Thr	Ile	Gly	Lys	Pro 265	Ile	Pro	Pro	Pro	Pro 270	Pro	Pro
Ser P	he Pro 27!		Pro	Pro	Pro	Pro 280	Pro	Gly	Thr	Gln	Leu 285	Pro	Pro	Pro
	ro Sei 90	r Tyr	Pro	Ser	Pro 295	Lys	Pro	Pro	Val	Gly 300	Pro	Gln	Ala	Ala
Asp I.	le Ty	r Met	Gln	Thr 310	Lys	Asn	Lys	Leu	Arg 315	His	Val	Glu	Thr	Glu 320
Ala L	eu Ly:	s Lys	Glu 325	Pro	Ser	Ser	Cys	Asp 330	Gly	His	Asp	Gly	Leu 335	Arg
Arg G	ln Ası	9 Ser 340	Ser	Arg	Lys	Pro	Arg 345	Ala	Phe	Ser	Lys	Gln 350	Pro	Ser
Thr G	ly As ₁ 35!		Tyr	Arg	Gln	Leu 360	Gly	Arg	Cys	Pro	Gly 365	Glu	Thr	Leu
Val A	la Aro	g Pro	Gly	Met	Ala 375	His	Arg	Glu	Glu	Ala 380	Glu	Leu	Pro	Gly
Asn H 385	is Val	l Pro	Asn	Gly 390	Cys	Ala	Ala	Asp	Pro 395	Lys	Ala	Ser	Arg	Glu 400
Gln G	ln Le	ı Pro	Pro 405	Pro	Pro	Pro	Pro	Pro 410	Pro	Leu	Pro	Glu	Ala 415	Ala
Ser S	er Pro	Pro 420	Pro	Val	Pro	Pro	Leu 425	Pro	Leu	Glu	Gly	Ala 430	Gly	Pro
Gly C	ys Gly 43!		Arg	Arg	Ser	Ser 440	Ser	Pro	Thr	Gly	Ser 445	Thr	Lys	Ser
	sn Val	l Met	Phe	Pro	Met 455	Gly	Asp	Asn	Ser	Glu 460	Leu	Leu	Ala	Glu
Ile L 465	ys Ala	a Gly	Lys	Ser 470	Leu	Lys	Pro	Thr	Pro 475	Gln	Ser	Lys	Gly	Leu 480
Thr T	hr Va	l Phe	Ser 485	Gly	Ser	Arg	Gln	Pro 490	Ala	Phe	Gln	Pro	Asp 495	Trp
Pro L	eu Pro	500	Val	Ser	Pro	Ala	Leu 505	Leu	Pro	Val	Arg	Ser 510	Pro	Thr

Pro Pro Ala Ala Gly Phe Gln Pro Leu Leu Asn Gly Ser Leu Val Pro Val Pro Pro Thr Thr Pro Ala Pro Gly Val Gln Leu Asp Val Glu Ala Leu Ile Pro Thr His Asp Glu Gln Gly Arg Pro Lys Pro Glu Trp Lys Arg Gln Val Met Val Gly Lys Met Gln Leu Lys Met Glu Glu Glu Glu Gln Arg Trp Lys Gln Arg Ala Ala Thr Gly Arg Ala Pro Arg Gln Arg Pro Lys Trp Thr Leu Pro Arg Ala Trp Ser Gly Gly Ser Gly Arg Ser Leu Thr Pro Ala Ser Pro Pro Ala Gly Gln Thr Arg Ser Leu Pro Ala Asp Ala Ala Pro Arg Ser His Tyr Thr Thr Gln Asp Met Gln Lys Leu Thr Ala Ala Ser Ser Cys Cys Tyr Pro Arg Glu Gly Trp Arg Tyr Pro Arg Glu Gly Trp Arg Tyr Ser Arg Glu His Asn Ala Ile Leu Trp Pro Phe Gly Glu Leu Met Thr Glu Ala Asp Ile Leu Arg Ile Glu Gln Gln Ser Arg Thr Cys Ser Cys Arg Pro Leu Thr Arg Ala Ser Arg Trp Arg Arg Cys Leu Arg Arg Pro Asp Cys Arg Gly Arg Phe Ala Trp Ala Ala Arg Thr Gly Ser Thr Gly Ala Ala Arg Leu Trp Arg Ala Arg Ser Ser Ser Ala Ala Ser Pro Cys Ser Ile Thr Ala Pro Pro Thr Ser Cys Ala His Trp Thr Arg Arg Pro Arg Ala Val Arg Ala Ala Ser Pro Arg

Ser Pro Leu 770	Ala Pro	Arg Ser 775	Ala S	Ser Pro		Cys A 780	Arg A	Arg	Thr	Thr
Trp Arg Pro	Ala Leu	Ala Ser 790	Pro A	Ala Pro	Pro E 795	Pro E	?ro I	Thr .	Ala	Arg 800
Trp Pro Thr	Gly Ser 805	Pro Trp	Thr F	Pro Trp 810	Ala A	Arg I	eu A		His 815	Arg
Ile Ala Arg	Arg Arg 820	Tyr Leu		Pro Ser 325	Ser 1	Frp A		Ala 330	Gly	Arg
Pro Ser Ala 835	-	Cys Ala	Ala S	Ser Arg	Thr 1		Ser I 345	Thr	Cys	Ala
Arg Ser Ala 850	Ser Phe	Thr Ser 855	Ser I	Irp Ser		Gly <i>A</i> 860	Ala S	Ser	Gly	Pro
Ser Ser Asp 865	Arg Ala	Phe Arg 870	Gly E	Pro Gly	Ala I 875	Pro A	Arg G	Gln	Thr	Ala 880
Pro Trp Arg	Asp Gly 885	Arg Pro	Cys 1	Pro Pro	Glu I	Leu G	Glu A		Thr 895	Asp
Ala Pro Arg	Leu Pro 900	Val Ser	_	Gly Glu 905	Ala H	His S		Pro 910	Asn	Glu
Arg Leu Arg 915		Leu Arg	Gln <i>F</i> 920	Arg Gln	Ala V		Gly I 925	Lys	Leu	Leu
His His Trp	Arg Ser	Leu Arg 935	Arg H	His Val		Pro S 940	Ser E	Pro	Gly	Leu
Ala His Gly 945	Val Tyr	Trp Pro 950	Gln H	His Phe	Leu 5 955	Ser E	Pro I	Leu	Asp	Gly 960
Gly Ala Pro	Pro Arg 965	Tyr Glu	Ser I	Leu Thr 970	Leu A	Asp I	Leu E		Met 975	Leu
Gly Tyr Phe	Gln Leu 980	Pro Glu		Gly Leu 985	Ser A	Arg G		Asp 990	Arg	Lys
Phe Arg His			Glu M 1000	Met Phe	His A	_	Leu <i>F</i> 005	Asp	Ser	His
Pro Trp Glu 1010	Arg Ile	Arg Leu 1015	Phe F	His Arg		Val I 020	Leu G	Glu	Glu	Val

Glu Ala Gly Arg Arg Gly Trp Ser Asp Gly Phe Glu Asp Leu Arg His 1025 1030 1035 1040

Arg Phe Phe Gly Asn Gly Leu Glu Ala Gly Pro Ala Pro Glu Glu Gln 1045 1050 1055

Ala Lys Lys Glu Glu Lys Gly Lys Glu Gln Glu Arg Thr Glu Glu
1060 1065 1070

Ala Ala Pro Val Gln Lys Gly Asp Pro Pro Lys Gly Gln Arg Glu Ala 1075 1080 1085

Leu Ala Pro Val Pro Gln Pro Pro Pro Pro Pro Ala Arg Pro Pro Ala 1090 1095 1100

Arg Arg Ala Ser Pro Pro Arg Leu Pro Gly Ser Gln Thr Leu Arg Val 1105 1110 1115 1120

Pro Lys Pro Pro Pro Lys Thr Leu Trp Asn 1125 1130

<210> 118

<211> 711

<212> PRT

<213> Homo sapiens

<400> 118

Met Gly Trp Leu Pro Leu Leu Leu Leu Thr Gln Cys Leu Gly Val 1 5 10 15

Pro Gly Gln Arg Ser Pro Leu Asn Asp Phe Gln Val Leu Arg Gly Thr 20 25 30

Glu Leu Gln His Leu Leu His Ala Val Val Pro Gly Pro Trp Gln Glu 35 40 45

Asp Val Ala Asp Ala Glu Glu Cys Ala Gly Arg Cys Gly Pro Leu Met 50 55 60

Asp Cys Arg Ala Phe His Tyr Asn Val Ser Ser His Gly Cys Gln Leu 65 70 75 80

Leu Pro Trp Thr Gln His Ser Pro His Thr Arg Leu Arg Arg Ser Gly 85 90 95

Arg Cys Asp Leu Phe Gln Lys Lys Asp Tyr Val Arg Thr Cys Ile Met
100 105 110

Asn A		Gly 115	Val	Gly	Tyr	Arg	Gly 120	Thr	Met	Ala	Thr	Thr 125	Val	Gly	Gly
Leu E	Pro 130	Cys	Gln	Ala	Trp	Ser 135	His	Lys	Phe	Pro	Asn 140	Asp	His	Lys	Tyr
Thr E	Pro	Thr	Leu	Arg	Asn 150	Gly	Leu	Glu	Glu	Asn 155	Phe	Cys	Arg	Asn	Pro 160
Asp G	Gly	Asp	Pro	Gly 165	Gly	Pro	Trp	Cys	Tyr 170	Thr	Thr	Asp	Pro	Ala 175	Val
Arg E	Phe	Gln	Ser 180	Cys	Gly	Ile	Lys	Ser 185	Cys	Arg	Glu	Ala	Ala 190	Cys	Val
Trp (-	Asn 195	Gly	Glu	Glu	Tyr	Arg 200	Gly	Ala	Val	Asp	Arg 205	Thr	Glu	Ser
Gly A	Arg 210	Glu	Cys	Gln	Arg	Trp 215	Asp	Leu	Gln	His	Pro 220	His	Gln	His	Pro
Phe 6	Glu	Pro	Gly	Lys	Phe 230	Leu	Asp	Gln	Gly	Leu 235	Asp	Asp	Asn	Tyr	Cys 240
Arg A	Asn	Pro	Asp	Gly 245	Ser	Glu	Arg	Pro	Trp 250	Cys	Tyr	Thr	Thr	Asp 255	Pro
Gln I	lle	Glu	Arg 260	Glu	Phe	Cys	Asp	Leu 265	Pro	Arg	Cys	Gly	Ser 270	Glu	Ala
Gln E		Arg 275	Gln	Glu	Ala	Thr	Thr 280	Val	Ser	Cys	Phe	Arg 285	Gly	Lys	Gly
Glu G	Gly 290	Tyr	Arg	Gly	Thr	Ala 295	Asn	Thr	Thr	Thr	Ala 300	Gly	Val	Pro	Cys
Gln <i>A</i> 305	Arg	Trp	Asp	Ala	Gln 310	Ile	Pro	His	Gln	His 315	Arg	Phe	Thr	Pro	Glu 320
Lys T	ſyr	Ala	Cys	Lys 325	Asp	Leu	Arg	Glu	Asn 330	Phe	Cys	Arg	Asn	Pro 335	Asp
Gly S	Ser	Glu	Ala 340	Pro	Trp	Cys	Phe	Thr 345	Leu	Arg	Pro	Gly	Met 350	Arg	Ala
Ala E		Cys 355	Tyr	Gln	Ile	Arg	Arg 360	Cys	Thr	Asp	Asp	Val 365	Arg	Pro	Gln

Asp	Cys 370	Tyr	His	Gly	Ala	Gly 375	Glu	Gln	Tyr	Arg	Gly 380	Thr	Val	Ser	Lys
Thr 385	Arg	Lys	Gly	Val	Gln 390	Cys	Gln	Arg	Trp	Ser 395	Ala	Glu	Thr	Pro	His 400
Lys	Pro	Gln	Phe	Thr 405	Phe	Thr	Ser	Glu	Pro 410	His	Ala	Gln	Leu	Glu 415	Glu
Asn	Phe	Cys	Arg 420	Asn	Pro	Asp	Gly	Asp 425	Ser	His	Gly	Pro	Trp 430	Cys	Tyr
Thr	Met	Asp 435	Pro	Arg	Thr	Pro	Phe 440	Asp	Tyr	Cys	Ala	Leu 445	Arg	Arg	Cys
Ala	Asp 450	Asp	Gln	Pro	Pro	Ser 455	Ile	Leu	Asp	Pro	Pro 460	Asp	Gln	Val	Gln
Phe 465	Glu	Lys	Cys	Gly	Lys 470	Arg	Val	Asp	Arg	Leu 475	Asp	Gln	Arg	Arg	Ser 480
Lys	Leu	Arg	Val	Val 485	Gly	Gly	His	Pro	Gly 490	Asn	Ser	Pro	Trp	Thr 495	Val
Ser	Leu	Arg	Asn 500	Arg	Gln	Gly	Gln	His 505	Phe	Cys	Gly	Gly	Ser 510	Leu	Val
Lys	Glu	Gln 515	Trp	Ile	Leu	Thr	Ala 520	Arg	Gln	Cys	Phe	Ser 525	Ser	Cys	His
Met	Pro 530	Leu	Thr	Gly	Tyr	Glu 535	Val	Trp	Leu	Gly	Thr 540	Leu	Phe	Gln	Asn
Pro 545	Gln	His	Gly	Glu	Pro 550	Ser	Leu	Gln	Arg	Val 555	Pro	Val	Ala	Lys	Met 560
Val	Cys	Gly	Pro	Ser 565	Gly	Ser	Gln	Leu	Val 570	Leu	Leu	Lys	Leu	Glu 575	Arg
Ser	Val	Thr	Leu 580	Asn	Gln	Arg	Val	Ala 585	Leu	Ile	Cys	Leu	Pro 590	Pro	Glu
Trp	Tyr	Val 595	Val	Pro	Pro	Gly	Thr 600	Lys	Cys	Glu	Ile	Ala 605	Gly	Trp	Gly
Glu	Thr 610	Lys	Gly	Thr	Gly	Asn 615	Asp	Thr	Val	Leu	Asn 620	Val	Ala	Phe	Leu

Asn Val Ile Ser Asn Gln Glu Cys Asn Ile Lys His Arg Gly Arg Val 625 630 635 640

Arg Glu Ser Glu Met Cys Thr Glu Gly Leu Leu Ala Pro Val Gly Ala 645 650 655

Cys Glu Gly Asp Tyr Gly Gly Pro Leu Ala Cys Phe Thr His Asn Cys 660 665 670

Trp Val Leu Glu Gly Ile Ile Ile Pro Asn Arg Val Cys Ala Arg Ser 675 680 685

Arg Trp Pro Ala Val Phe Thr Arg Val Ser Val Phe Val Asp Trp Ile 690 695 700

His Lys Val Met Arg Leu Gly 705 710

<210> 119

<211> 711

<212> PRT

<213> Homo sapiens

<400> 119

Met Gly Trp Leu Pro Leu Leu Leu Leu Thr Gln Cys Leu Gly Val
1 5 10 15

Pro Gly Gln Arg Ser Pro Leu Asn Asp Phe Gln Val Leu Arg Gly Thr 20 25 30

Glu Leu Gln His Leu Leu His Ala Val Val Pro Gly Pro Trp Gln Glu 35 40 45

Asp Val Ala Asp Ala Glu Glu Cys Ala Gly Arg Cys Gly Pro Leu Met
50 55 60

Asp Cys Arg Ala Phe His Tyr Asn Val Ser Ser His Gly Cys Gln Leu 65 70 75 80

Leu Pro Trp Thr Gln His Ser Pro His Thr Arg Leu Arg Arg Ser Gly
85 90 95

Arg Cys Asp Leu Phe Gln Lys Lys Asp Tyr Val Arg Thr Cys Ile Met 100 105 110

Asn Asn Gly Val Gly Tyr Arg Gly Thr Met Ala Thr Thr Val Gly Gly

115	120	125

Leu	Pro 130	Cys	Gln	Ala	Trp	Ser 135	His	Lys	Phe	Pro	Asn 140	Asp	His	Lys	Tyr
Thr 145	Pro	Thr	Leu	Arg	Asn 150	Gly	Leu	Glu	Glu	Asn 155	Phe	Cys	Arg	Asn	Pro 160
Asp	Gly	Asp	Pro	Gly 165	Gly	Pro	Trp	Cys	Tyr 170	Thr	Thr	Asp	Pro	Ala 175	Val
Arg	Phe	Gln	Ser 180	Cys	Gly	Ile	Lys	Ser 185	Cys	Arg	Glu	Ala	Ala 190	Cys	Val
Trp	Cys	Asn 195	Gly	Glu	Glu	Tyr	Arg 200	Gly	Ala	Val	Asp	Arg 205	Thr	Glu	Ser
Gly	Arg 210	Glu	Cys	Gln	Arg	Trp 215	Asp	Leu	Gln	His	Pro 220	His	Gln	His	Pro
Phe 225	Glu	Pro	Gly	Lys	Phe 230	Leu	Asp	Gln	Gly	Leu 235	Asp	Asp	Asn	Tyr	Cys 240
Arg	Asn	Pro	Asp	Gly 245	Ser	Glu	Arg	Pro	Trp 250	Cys	Tyr	Thr	Thr	Asp 255	Pro
Gln	Ile	Glu	Arg 260	Glu	Phe	Cys	Asp	Leu 265	Pro	Arg	Cys	Gly	Ser 270	Glu	Ala
Gln	Pro	Arg 275	Gln	Glu	Ala	Thr	Thr 280	Val	Ser	Cys	Phe	Arg 285	Gly	Lys	Gly
Glu	Gly 290	Tyr	Arg	Gly	Thr	Ala 295	Asn	Thr	Thr	Thr	Ala 300	Gly	Val	Pro	Cys
Gln 305	Arg	Trp	Asp	Ala	Gln 310	Ile	Pro	His	Gln	His 315	Arg	Phe	Thr	Pro	Glu 320
Lys	Tyr	Ala	Cys	Lys 325	Asp	Leu	Arg	Glu	Asn 330	Phe	Cys	Arg	Asn	Pro 335	Asp
Gly	Ser	Glu	Ala 340	Pro	Trp	Cys	Phe	Thr 345	Leu	Arg	Pro	Gly	Met 350	Arg	Ala
Ala	Phe	Cys 355	Tyr	Gln	Ile	Arg	Arg 360	Cys	Thr	Asp	Asp	Val 365	Arg	Pro	Gln

Asp Cys Tyr His Gly Ala Gly Glu Gln Tyr Arg Gly Thr Val Ser Lys

370	375	380

Thr 385	Arg	Lys	Gly	Val	Gln 390	Cys	Gln	Arg	Trp	Ser 395	Ala	Glu	Thr	Pro	His 400
Lys	Pro	Gln	Phe	Thr 405	Phe	Thr	Ser	Glu	Pro 410	His	Ala	Gln	Leu	Glu 415	Glu
Asn	Phe	Cys	Arg 420	Asn	Pro	Asp	Gly	Asp 425	Ser	His	Gly	Pro	Trp 430	Cys	Tyr
Thr	Met	Asp 435	Pro	Arg	Thr	Pro	Phe 440	Asp	Tyr	Cys	Ala	Leu 445	Arg	Arg	Cys
Ala	Asp 450	Asp	Gln	Pro	Pro	Ser 455	Ile	Leu	Asp	Pro	Pro 460	Asp	Gln	Val	Gln
Phe 465	Glu	Lys	Cys	Gly	Lys 470	Arg	Val	Asp	Arg	Leu 475	Asp	Gln	Arg	Arg	Ser 480
Lys	Leu	Arg	Val	Val 485	Gly	Gly	His	Pro	Gly 490	Asn	Ser	Pro	Trp	Thr 495	Val
Ser	Leu	Arg	Asn 500	Arg	Gln	Gly	Gln	His 505	Phe	Cys	Gly	Gly	Ser 510	Leu	Val
Lys	Glu	Gln 515	Trp	Ile	Leu	Thr	Ala 520	Arg	Gln	Cys	Phe	Ser 525	Ser	Cys	His
Met	Pro 530	Leu	Thr	Gly	Tyr	Glu 535	Val	Trp	Leu	Gly	Thr 540	Leu	Phe	Gln	Asn
Pro 545	Gln	His	Gly	Glu	Pro 550	Ser	Leu	Gln	Arg	Val 555	Pro	Val	Ala	Lys	Met 560
Val	Cys	Gly	Pro	Ser 565	Gly	Ser	Gln	Leu	Val 570	Leu	Leu	Lys	Leu	Glu 575	Arg
Ser	Val	Thr	Leu 580	Asn	Gln	Arg	Val	Ala 585	Leu	Ile	Cys	Leu	Pro 590	Pro	Glu
Trp	Tyr	Val 595	Val	Pro	Pro	Gly	Thr 600	Lys	Cys	Glu	Ile	Ala 605	Gly	Trp	Gly
Glu	Thr 610	Lys	Gly	Thr	Gly	Asn 615	Asp	Thr	Val	Leu	Asn 620	۷al	Ala	Leu	Leu

Asn Val Ile Ser Asn Gln Glu Cys Asn Ile Lys His Arg Gly Arg Val

Arg Glu Ser Glu Met Cys Thr Glu Gly Leu Leu Ala Pro Val Gly Ala 645 650 655

Cys Glu Gly Asp Tyr Gly Gly Pro Leu Ala Cys Phe Thr His Asn Cys 660 665 670

Trp Val Leu Glu Gly Ile Ile Ile Pro Asn Arg Val Cys Ala Arg Ser 675 680 685

Arg Trp Pro Ala Val Phe Thr Arg Val Ser Val Phe Val Asp Trp Ile 690 695 700

His Lys Val Met Arg Leu Gly 705 710

<210> 120

<211> 711

<212> PRT

<213> Homo sapiens

<400> 120

Met Gly Trp Leu Pro Leu Leu Leu Leu Thr Gln Tyr Leu Gly Val 1 5 10 15

Pro Gly Gln Arg Ser Pro Leu Asn Asp Phe Gln Val Leu Arg Gly Thr 20 25 30

Glu Leu Gln His Leu Leu His Ala Val Val Pro Gly Pro Trp Gln Glu 35 40 45

Asp Val Ala Asp Ala Glu Glu Cys Ala Gly Arg Cys Gly Pro Leu Met 50 55 60

Asp Cys Arg Ala Phe His Tyr Asn Val Ser Ser His Gly Cys Gln Leu 65 70 75 80

Leu Pro Trp Thr Gln His Ser Pro His Thr Arg Leu Arg Arg Ser Gly 85 90 95

Arg Cys Asp Leu Phe Gln Lys Lys Asp Tyr Val Arg Thr Cys Ile Met
100 105 110

Asn Asn Gly Val Gly Tyr Arg Gly Thr Met Ala Thr Thr Val Gly Gly 115 120 125

Leu Pro Cys Gln Ala Trp Ser His Lys Phe Pro Asn Asp His Lys Tyr Thr Pro Thr Leu Arg Asn Gly Leu Glu Glu Asn Phe Cys Arg Asn Pro Asp Gly Asp Pro Gly Gly Pro Trp Cys Tyr Thr Thr Asp Pro Ala Val Arg Phe Gln Ser Cys Gly Ile Lys Ser Cys Arg Glu Ala Ala Cys Val Trp Cys Asn Gly Glu Glu Tyr Arg Gly Ala Val Asp Arg Thr Glu Ser Gly Arg Glu Cys Gln Arg Trp Asp Leu Gln His Pro His Gln His Pro Phe Glu Pro Gly Lys Phe Leu Asp Gln Gly Leu Asp Asp Asn Tyr Cys Arg Asn Pro Asp Gly Ser Glu Arg Pro Trp Cys Tyr Thr Thr Asp Pro Gln Ile Glu Arg Glu Phe Cys Asp Leu Pro Arg Cys Gly Ser Glu Ala Gln Pro Arg Gln Glu Ala Thr Thr Val Ser Cys Phe Arg Gly Lys Gly Glu Gly Tyr Arg Gly Thr Ala Asn Thr Thr Thr Ala Gly Val Pro Cys Gln Arg Trp Asp Ala Gln Ile Pro His Gln His Arg Phe Thr Pro Glu Lys Tyr Ala Cys Lys Asp Leu Arg Glu Asn Phe Cys Arg Asn Pro Asp Gly Ser Glu Ala Pro Trp Cys Phe Thr Leu Arg Pro Gly Met Arg Ala Ala Phe Cys Tyr Gln Ile Arg Arg Cys Thr Asp Asp Val Arg Pro Gln Asp Cys Tyr His Gly Ala Gly Glu Gln Tyr Arg Gly Thr Val Ser Lys

Thr 385	Arg	Lys	Gly	Val	Gln 390	Cys	Gln	Arg	Trp	Ser 395	Ala	Glu	Thr	Pro	His 400
Lys	Pro	Gln	Phe	Thr 405	Phe	Thr	Ser	Glu	Pro 410	His	Ala	Gln	Leu	Glu 415	Glu
Asn	Phe	Cys	Arg 420	Asn	Pro	Asp	Gly	Asp 425	Ser	His	Gly	Pro	Trp 430	Cys	Tyr
Thr	Met	Asp 435	Pro	Arg	Thr	Pro	Phe 440	Asp	Tyr	Cys	Ala	Leu 445	Arg	Arg	Cys
Ala	Asp 450	Asp	Gln	Pro	Pro	Ser 455	Ile	Leu	Asp	Pro	Pro 460	Asp	Gln	Val	Gln
Phe 465	Glu	Lys	Cys	Gly	Lys 470	Arg	Val	Asp	Arg	Leu 475	Asp	Gln	Arg	Arg	Ser 480
Lys	Leu	Arg	Val	Val 485	Gly	Gly	His	Pro	Gly 490	Asn	Ser	Pro	Trp	Thr 495	Val
Ser	Leu	Arg	Asn 500	Arg	Gln	Gly	Gln	His 505	Phe	Cys	Gly	Gly	Ser 510	Leu	Val
Lys	Glu	Gln 515	Trp	Ile	Leu	Thr	Ala 520	Arg	Gln	Cys	Phe	Ser 525	Ser	Cys	His
Met	Pro 530	Leu	Thr	Gly	Tyr	Glu 535	Val	Trp	Leu	Gly	Thr 540	Leu	Phe	Gln	Asn
Pro 545	Gln	His	Gly	Glu	Pro 550	Ser	Leu	Gln	Arg	Val 555	Pro	Val	Ala	Lys	Met 560
Val	Cys	Gly	Pro	Ser 565	Gly	Ser	Gln	Leu	Val 570	Leu	Leu	Lys	Leu	Glu 575	Arg
Ser	Val	Thr	Leu 580	Asn	Gln	Arg	Val	Ala 585	Leu	Ile	Cys	Leu	Pro 590	Pro	Glu
Trp	Tyr	Val 595	Val	Pro	Pro	Gly	Thr 600	Lys	Cys	Glu	Ile	Ala 605	Gly	Trp	Gly
Glu	Thr 610	Lys	Gly	Thr	Gly	Asn 615	Asp	Thr	Val	Leu	Asn 620	Val	Ala	Leu	Leu
Asn 625	Val	Ile	Ser	Asn	Gln 630	Glu	Cys	Asn	Ile	Lys 635	His	Arg	Gly	Arg	Val 640

Arg Glu Ser Glu Met Cys Thr Glu Gly Leu Leu Ala Pro Val Gly Ala 645 650 655

Cys Glu Gly Asp Tyr Gly Gly Pro Leu Ala Cys Phe Thr His Asn Cys 660 665 670

Trp Val Leu Glu Gly Ile Ile Ile Pro Asn Arg Val Cys Ala Arg Ser 675 680 685

Arg Trp Pro Ala Val Phe Thr Arg Val Ser Val Phe Val Asp Trp Ile 690 695 700

His Lys Val Met Arg Leu Gly 705 710

<210> 121

<211> 567

<212> PRT

<213> Homo sapiens

<400> 121

Met Thr Ser Arg Cys Ser Gly Ala Gln Ser Tyr Leu Leu His Ala Val 1 5 10 15

Val Pro Gly Pro Trp Gln Glu Asp Val Ala Asp Ala Glu Glu Cys Ala 20 25 30

Gly Arg Cys Gly Leu Leu Met Asp Cys Trp Ala Phe His Tyr Asn Val 35 40 45

Ser Ser His Gly Cys Gln Leu Leu Pro Trp Thr Gln His Ser Pro His 50 55 60

Ser Arg Leu Arg His Ser Gly Arg Cys Asp Leu Phe Gln Lys Lys Asp 65 70 75 80

Tyr Ile Arg Thr Cys Ile Met Asn Asn Gly Val Gly Tyr Arg Asp Thr 85 90 95

Met Ala Thr Thr Val Gly Gly Leu Ser Cys Gln Ala Trp Ser His Lys
100 105 110

Phe Pro Asn Asp His Gln Tyr Met Pro Thr Leu Arg Asn Gly Leu Glu 115 120 125

Glu Asn Phe Cys Arg Asn Pro Asp Gly Asp Pro Gly Gly Pro Trp Cys
130 135 140

His 145	Thr	Thr	Asp	Pro	Ala 150	Val	Arg	Phe	Gln	Ser 155	Cys	Gly	Ile	Lys	Ser 160
Cys	Leu	Val	Ala	Ala 165	Cys	Val	Trp	Суз	Asn 170	Gly	Glu	Glu	Tyr	Arg 175	Gly
Ala	Val	Asp	Arg 180	Thr	Glu	Ser	Gly	Arg 185	Glu	Cys	Gln	Arg	Trp 190	Asp	Leu
Gln	His	Pro 195	His	Gln	His	Pro	Phe 200	Glu	Pro	Gly	Lys	Phe 205	Leu	Asp	Gln
Gly	Leu 210	Asp	Asp	Asn	Tyr	Cys 215	Arg	Ser	Pro	Asp	Gly 220	Ser	Gln	Arg	Pro
Trp 225	Cys	Tyr	Thr	Thr	Asp 230	Pro	Gln	Ile	Glu	Arg 235	Glu	Phe	Cys	Asp	Leu 240
Pro	Arg	Cys	Gly	Ser 245	Glu	Ala	Gln	Pro	Arg 250	Gln	Glu	Ala	Thr	Ser 255	Val
Ser	Cys	Phe	Arg 260	Gly	Lys	Gly	Glu	Gly 265	Tyr	Arg	Gly	Thr	Ala 270	Asn	Thr
Thr	Thr	Ala 275	Gly	Val	Pro	Cys	Gln 280	Arg	Trp	Asp	Ala	Gln 285	Ile	Pro	His
Gln	His 290	Arg	Phe	Thr	Pro	Glu 295	Lys	Tyr	Ala	Cys	Lys 300	Asp	Leu	Arg	Glu
Asn 305	Phe	Cys	Arg	Asn	Pro 310	Asp	Gly	Ser	Glu	Ala 315	Pro	Trp	Cys	Phe	Thr 320
Leu	Arg	Pro	Gly	Thr 325	Arg	Val	Gly	Phe	Cys 330	Tyr	Gln	Ile	Arg	Arg 335	Cys
Thr	Asp	Asp	Val 340	Arg	Pro	Gln	Asp	Cys 345	Tyr	His	Gly	Ala	Gly 350	Glu	Gln
Tyr	Arg	Gly 355	Thr	Val	Ser	Lys	Thr 360	Arg	Lys	Gly	Val	Gln 365	Cys	Gln	Arg
Trp	Ser 370	Ala	Glu	Thr	Pro	His 375	Lys	Pro	Gln	Phe	Thr 380	Phe	Thr	Ser	Glu
Pro 385	His	Ala	Gln	Leu	Glu 390	Glu	Asn	Phe	Cys	Gln 395	Asn	Pro	Asp	Gly	Asp 400

Ser His Gly Pro Trp Cys Tyr Thr Met Asp Pro Arg Thr Pro Phe Asp 405 410 415 Tyr Cys Ala Leu Arg Arg Cys Ala Asp Asp Gln Pro Pro Ser Ile Leu 430 420 425 Asp Pro Pro Asp Gln Val Gln Phe Glu Lys Cys Gly Lys Arg Val Asp 440 Arg Leu Asp Gln Arg Arg Ser Lys Leu Arg Val Ala Gly Gly His Pro 455 Gly Asn Ser Pro Trp Thr Val Ser Leu Gly Asn Arg Gln Gly Gln His 465 470 475 Phe Cys Gly Gly Ser Leu Val Lys Glu Gln Trp Ile Leu Thr Ala Arg 485 490 495 Gln Cys Phe Ser Ser Cys His Met Pro Leu Thr Gly Tyr Glu Val Trp 505 500 Leu Gly Thr Leu Phe Gln Asn Pro Gln His Gly Glu Pro Gly Leu Gln 520 525 515 Arg Val Pro Val Ala Lys Met Leu Cys Gly Pro Ser Gly Ser Gln Leu 535 Val Leu Leu Lys Leu Glu Arg Ser Val Thr Leu Asn Gln Arg Val Ala 545 550 555 Leu Ile Cys Leu Pro Pro Glu 565 <210> 122 <211> 78 <212> PRT <213> Artificial Sequence <220>

<223> Description of Artificial Sequence: Kringle domain sequence

<400> 122

Cys Val Trp Cys Asn Gly Glu Glu Tyr Arg Gly Ala Val Asp Arg Thr 1 10 15

Glu Ser Gly Arg Glu Cys Gln Arg Trp Asp Leu Gln His Pro His Gln 20 25 30

His Pro Phe Glu Pro Gly Lys Phe Leu Asp Gln Gly Leu Asp Asp Asn 35 40 45

Tyr Cys Arg Asn Pro Asp Gly Ser Glu Arg Pro Trp Cys Tyr Thr Thr 50 55 60

Asp Pro Gln Ile Glu Arg Glu Phe Cys Asp Leu Pro Arg Cys 65 70 75

<210> 123

<211> 79

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Kringle domain sequence

<400> 123

Cys Phe Arg Gly Lys Gly Glu Gly Tyr Arg Gly Thr Ala Asn Thr Thr

1 5 10 15

Thr Ala Gly Val Pro Cys Gln Arg Trp Asp Ala Gln Ile Pro His Gln 20 25 30

His Arg Phe Thr Pro Glu Lys Tyr Ala Cys Lys Asp Leu Arg Glu Asn 35 40 45

Phe Cys Arg Asn Pro Asp Gly Ser Glu Ala Pro Trp Cys Phe Thr Leu 50 55 60

Arg Pro Gly Met Arg Ala Ala Phe Cys Tyr Gln Ile Arg Arg Cys 65 70 75

<210> 124

<211> 77

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Kringle domain sequence

<400> 124

Cys Ile Met Asn Asn Gly Val Gly Tyr Arg Gly Thr Met Ala Thr Thr

1 5 10 15

Val Gly Gly Leu Pro Cys Gln Ala Trp Ser His Lys Phe Pro Asn Asp 20 25 30

His Lys Tyr Thr Pro Thr Leu Arg Asn Gly Leu Glu Glu Asn Phe Cys 35 40 45

Arg Asn Pro Asp Gly Asp Pro Gly Gly Pro Trp Cys Tyr Thr Thr Asp 50 55 60

Pro Ala Val Arg Phe Gln Ser Cys Gly Ile Lys Ser Cys 65 70 75

<210> 125

<211> 80

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Kringle domain sequence

<400> 125

Cys Val Trp Cys Asn Gly Glu Glu Tyr Arg Gly Ala Val Asp Arg Thr 1 5 10 15

Glu Ser Gly Arg Glu Cys Gln Arg Trp Asp Leu Gln His Pro His Gln
20 25 30

His Pro Phe Glu Pro Gly Lys Phe Leu Asp Gln Gly Leu Asp Asp Asn 35 40 45

Tyr Cys Arg Asn Pro Asp Gly Ser Glu Arg Pro Trp Cys Tyr Thr Thr 50 55 60

Asp Pro Gln Ile Glu Arg Glu Phe Cys Asp Leu Pro Arg Cys Gly Ser 65 70 75 80

<210> 126

<211> 79

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Kringle domain sequence

<400> 126

Arg Thr Cys Ile Met Asn Asn Gly Val Gly Tyr Arg Gly Thr Met Ala $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Thr Thr Val Gly Gly Leu Pro Cys Gln Ala Trp Ser His Lys Phe Pro 20 25 30

Asn Asp His Lys Tyr Thr Pro Thr Leu Arg Asn Gly Leu Glu Glu Asn 35 40 45

Phe Cys Arg Asn Pro Asp Gly Asp Pro Gly Gly Pro Trp Cys Tyr Thr 50 55 60

Thr Asp Pro Ala Val Arg Phe Gln Ser Cys Gly Ile Lys Ser Cys 65 70 75

<210> 127

<211> 81

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Kringle domain sequence

<400> 127

Cys Phe Arg Gly Lys Gly Glu Gly Tyr Arg Gly Thr Ala Asn Thr Thr 1 5 10 15

Thr Ala Gly Val Pro Cys Gln Arg Trp Asp Ala Gln Ile Pro His Gln 20 25 30

His Arg Phe Thr Pro Glu Lys Tyr Ala Cys Lys Asp Leu Arg Glu Asn 35 40 45

Phe Cys Arg Asn Pro Asp Gly Ser Glu Ala Pro Trp Cys Phe Thr Leu 50 55 60

Arg Pro Gly Met Arg Ala Ala Phe Cys Tyr Gln Ile Arg Arg Cys Thr 65 70 75 80

Asp

<210> 128

<211> 80

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Apple domain sequence

<400> 128

Asp Phe Gln Val Leu Arg Gly Thr Glu Leu Gln His Leu Leu His Ala 1 5 10 15

Val Val Pro Gly Pro Trp Gln Glu Asp Val Ala Asp Ala Glu Glu Cys
20 25 30

Ala Gly Arg Cys Gly Pro Leu Met Asp Cys Arg Ala Phe His Tyr Asn 35 40 45

Val Ser Ser His Gly Cys Gln Leu Leu Pro Trp Thr Gln His Ser Pro 50 55 60

His Thr Arg Leu Arg Arg Ser Gly Arg Cys Asp Leu Phe Gln Lys Lys 65 70 75 80

<210> 129

<211> 431

<212> PRT

<213> Mus musculus

<400> 129

Met Asp Ala Arg Trp Trp Ala Val Val Leu Ala Thr Leu Pro Ser 1 5 10 15

Leu Gly Ala Gly Glu Ser Pro Glu Ala Pro Pro Gln Ser Trp Thr
20 25 30

Gln Leu Trp Leu Phe Arg Phe Leu Leu Asn Val Ala Gly Tyr Ala Ser 35 40 45 Phe Met Val Pro Gly Tyr Leu Leu Val Gln Tyr Leu Arg Arg Lys Asn Tyr Leu Glu Thr Gly Arg Gly Leu Cys Phe Pro Leu Val Lys Ala Cys Val Phe Gly Asn Glu Pro Lys Ala Pro Asp Glu Val Leu Leu Ala Pro Arg Thr Glu Thr Ala Glu Ser Thr Pro Ser Trp Gln Val Leu Lys Leu Val Phe Cys Ala Ser Gly Leu Gln Val Ser Tyr Leu Thr Trp Gly Ile Leu Gln Glu Arg Val Met Thr Gly Ser Tyr Gly Ala Thr Ala Thr Ser Pro Gly Glu His Phe Thr Asp Ser Gln Phe Leu Val Leu Met Asn Arg Val Leu Ala Leu Val Val Ala Gly Leu Tyr Cys Val Leu Arg Lys Gln Pro Arq His Gly Ala Pro Met Tyr Arg Tyr Ser Phe Ala Ser Leu Ser Asn Val Leu Ser Ser Trp Cys Gln Tyr Glu Ala Leu Lys Phe Val Ser Phe Pro Thr Gln Val Leu Ala Lys Ala Ser Lys Val Ile Pro Val Met Met Met Gly Lys Leu Val Ser Arg Arg Ser Tyr Glu His Trp Glu Tyr Leu Thr Ala Gly Leu Ile Ser Ile Gly Val Ser Met Phe Leu Leu Ser Ser Gly Pro Glu Pro Arg Ser Ser Pro Ala Thr Thr Leu Ser Gly Leu Val Leu Leu Ala Gly Tyr Ile Ala Phe Asp Ser Phe Thr Ser Asn Trp Gln Asp Ala Leu Phe Ala Tyr Lys Met Ser Ser Val Gln Met Met Phe

Gly Val Asn Leu Phe Ser Cys Leu Phe Thr Val Gly Ser Leu Leu Glu Gln Gly Ala Leu Leu Glu Gly Ala Arg Phe Met Gly Arg His Ser Glu Phe Ala Leu His Ala Leu Leu Ser Ile Cys Ser Ala Phe Gly Gln Leu Phe Ile Phe Tyr Thr Ile Gly Gln Phe Gly Ala Ala Val Phe Thr Ile Ile Met Thr Leu Arg Gln Ala Ile Ala Ile Leu Leu Ser Cys Leu Leu Tyr Gly His Thr Val Thr Val Gly Gly Leu Gly Val Ala Val Val Phe Thr Ala Leu Leu Leu Arg Val Tyr Ala Arg Gly Arg Lys Gln Arg Gly Lys Lys Ala Val Pro Thr Glu Pro Pro Val Gln Lys Val <210> 130 <211> 465 <212> PRT <213> Drosophila melanogaster <400> 130 Met Tyr Ala Tyr Asn Lys Met Gly Arg Val Pro Glu Leu Val Ile Cys Ser Phe Ile Val Val Thr Leu Leu Val Ile His Phe Phe Ser Asp Leu Leu Arg Ala Ser Leu Gly Gly Tyr Tyr Asn Gln Asp Val Thr Leu Ser Gln Leu Val Glu Ser Gln Asn Ser Asp Tyr Ala Trp Phe Leu Lys Leu

Leu Val Asn Cys Phe Gly Tyr Ser Cys Val Phe Val Pro Gly Phe Leu

Ile Tyr Lys Tyr Val Gly Arg Ile Asn Tyr Leu Glu Arg Gly Asn Lys

Thr	Phe	Leu	His 100	Lys	Ala	Ile	Asn	Met 105	Cys	Ile	Thr	Gly	Asn 110	Ser	Gly
Tyr	Asp	Gln 115	Leu	Asp	Ala	Gly	Thr 120	Ser	Thr	Ala	Asp	Lys 125	Asp	Arg	Pro
Ala	Ala 130	Ser	Thr	Ala	Pro	Lys 135	Arg	Thr	Ser	Ser	Gln 140	Glu	Ala	Val	Gln
Leu 145	Leu	Trp	Cys	Phe	Gly 150	Gly	Leu	Met	Ile	Ser 155	Tyr	Leu	Thr	Trp	Gly 160
Val	Leu	Gln	Glu	Lys 165	Ile	Met	Thr	Gln	Asn 170	Tyr	Leu	Asn	Phe	Thr 175	Gly
Glu	Ser	Ala	Lys 180	Phe	Lys	Asp	Ser	Gln 185	Phe	Leu	Val	Phe	Ser 190	Asn	Arg
Leu	Leu	Ala 195	Phe	Leu	Val	Ala	Leu 200	Ala	Tyr	Leu	Gln	Trp 205	Gln	Pro	Ser
Pro	Val 210	Arg	His	Arg	Ala	Pro 215	Leu	Tyr	Lys	Tyr	Ser 220	Tyr	Ala	Ser	Phe
Ser 225	Asn	Ile	Met	Ser	Ala 230	Trp	Phe	Gln	Tyr	Glu 235	Ala	Leu	Lys	Phe	Val 240
Asn	Phe	Pro	Thr	Gln 245	Val	Leu	Ala	Lys	Ser 250	Cys	Lys	Ile	Ile	Pro 255	Val
Met	Leu	Met	Gly 260	Lys	Ile	Met	Ser	Lys 265	Ala	Lys	Tyr	Glu	Ser 270	Tyr	Glu
Tyr	Val	Thr 275	Ala	Leu	Leu	Ile	Ser 280	Leu	Gly	Met	Ile	Phe 285	Phe	Met	Ser
Gly	Ser 290	Ser	Asp	Ser	Ser	Lys 295	Ala	Ser	Gly	Val	Thr 300	Thr	Leu	Thr	Gly
Ile 305	Phe	Leu	Leu	Ser	Met 310	Tyr	Met	Val	Phe	Asp 315	Ser	Phe	Thr	Ala	Asn 320
Trp	Gln	Gly	Ser	Leu	Phe	Lys	Ser	Tyr	Gly	Met	Thr	Pro	Leu	Gln	Met

Met Cys Gly Val Asn Leu Phe Ser Ser Ile Phe Thr Gly Ala Ser Leu

340 345 350

Ser Met Gln Gly Gly Phe Met Asp Ser Leu Ala Phe Ala Thr Glu His 355 360 365

Pro Lys Phe Val Phe Asp Met Val Val Leu Ser Val Cys Ser Ala Val 370 380

Gly Gln Leu Phe Ile Tyr His Thr Ile Asp Val Phe Gly Pro Val Val 385 390 395 400

Phe Thr Ile Ile Met Thr Leu Arg Gln Ala Val Ala Ile Met Leu Ser 405 410 415

Cys Phe Ile Tyr Gln His Ser Ile Ser Leu Leu Gly Ile Phe Gly Val \$420\$ \$425\$ \$430

Leu Ile Val Phe Val Ala Ile Phe Leu Arg Val Tyr Cys Thr Gln Arg
435 440 445

Leu Arg Ala Ile Arg Lys Arg Ala Glu Ala Asn Lys Pro Lys Met Ala 450 455 460

Val 465

<210> 131

<211> 465

<212> PRT

<213> Drosophila melanogaster

<400> 131

Met Tyr Ala Tyr Asn Lys Met Gly Arg Val Pro Glu Leu Val Ile Cys 1 5 10 15

Ser Phe Ile Val Val Ser Leu Leu Val Ile His Phe Phe Ser Asp Leu 20 25 30

Leu Arg Ala Ser Leu Gly Gly Tyr Tyr Asn Gln Asp Val Thr Leu Ser
35 40 45

Gln Leu Val Glu Ser Gln Asn Ser Asp Tyr Ala Trp Phe Leu Lys Leu 50 55 60

Leu Val Asn Cys Phe Gly Tyr Ser Cys Val Phe Val Pro Gly Phe Leu 65 70 75 80

Ile Tyr Lys Tyr Val Gly Arg Ile Asn Tyr Leu Glu Arg Gly Asn Lys Thr Phe Leu His Lys Ala Ile Asn Met Cys Ile Thr Gly Asn Ser Gly Tyr Asp Gln Leu Asp Ala Gly Thr Ser Thr Ala Asp Lys Asp Arg Pro Ala Ala Ser Thr Ala Pro Lys Arg Thr Ser Ser Gln Glu Ala Val Gln Leu Leu Trp Cys Phe Gly Gly Leu Met Ile Ser Tyr Leu Thr Trp Gly Val Leu Gln Glu Lys Ile Met Thr Gln Asn Tyr Leu Asn Phe Thr Gly Glu Ser Ala Lys Phe Lys Asp Ser Gln Phe Leu Val Phe Ser Asn Arg Leu Leu Ala Phe Leu Val Ala Leu Ala Tyr Leu Gln Trp Gln Pro Ser Pro Val Arg His Arg Ala Pro Leu Tyr Lys Tyr Ser Tyr Ala Ser Phe Ser Asn Ile Met Ser Ala Trp Phe Gln Tyr Glu Ala Leu Lys Phe Val Asn Phe Pro Thr Gln Val Leu Ala Lys Ser Cys Lys Ile Ile Pro Val Met Leu Met Gly Lys Ile Met Ser Lys Ala Lys Tyr Glu Ser Tyr Glu Tyr Val Thr Ala Leu Leu Ile Ser Leu Gly Met Ile Phe Phe Met Ser Gly Ser Ser Asp Ser Ser Lys Ala Ser Gly Val Thr Thr Leu Thr Gly Ile Phe Leu Leu Ser Met Tyr Met Val Phe Asp Ser Phe Thr Ala Asn Trp Gln Gly Ser Leu Phe Lys Ser Tyr Gly Met Thr Pro Leu Gln Met

Met Cys Gly Val Asn Leu Phe Ser Ser Ile Phe Thr Gly Ala Ser Leu Ser Met Gln Gly Gly Phe Met Asp Ser Leu Ala Phe Ala Thr Glu His Pro Lys Phe Val Phe Asp Met Val Val Leu Ser Val Cys Ser Ala Val Gly Gln Leu Phe Ile Tyr His Thr Ile Asp Val Phe Gly Pro Val Val Phe Thr Ile Ile Met Thr Leu Arg Gln Ala Val Ala Ile Met Leu Ser Cys Phe Ile Tyr Gln His Ser Ile Ser Leu Leu Gly Ile Phe Gly Val Leu Ile Val Phe Val Ala Ile Phe Leu Arg Val Tyr Cys Thr Gln Arg Leu Arg Ala Ile Arg Lys Arg Ala Glu Ala Asn Lys Pro Lys Met Ala Val <210> 132 <211> 417 <212> PRT <213> Caenorhabditis elegans <400> 132 Met Asp Arg Ser Ile Met Pro Ile Asp Ser Pro Ala Arg Asp Lys Pro Pro Asp Glu Leu Val Trp Pro Leu Arg Leu Phe Leu Ile Leu Leu Gly Tyr Ser Thr Val Ala Thr Pro Ala Ala Ile Leu Ile Tyr Tyr Val Arg Arg Asn Arg His Ala Phe Glu Thr Pro Tyr Leu Ser Ile Arg Leu Leu

Leu Arg Ser Phe Ala Val Gly Asn Pro Glu Tyr Gln Leu Ile Pro Thr

Gly	Glu	Lys	Gln	Ala 85	Arg	Lys	Glu	Asn	Asp 90	Ser	Ile	Pro	Gln	Thr 95	Arg
Ala	Gln	Cys	Ile 100	Asn	Val	Ile	Ile	Leu 105	Leu	Leu	Phe	Phe	Phe 110	Ser	Gly
Ile	Gln	Val 115	Thr	Leu	Val	Ala	Met 120	Gly	Val	Leu	Gln	Glu 125	Arg	Ile	Ile
Thr	Arg 130	Gly	Tyr	Arg	Arg	Ser 135	Asp	Gln	Leu	Glu	Val 140	Glu	Asp	Lys	Phe
Gly 145	Glu	Thr	Gln	Phe	Leu 150	Ile	Phe	Cys	Asn	Arg 155	Ile	Val	Ala	Leu	Val 160
Leu	Ser	Leu	Met	Ile 165	Leu	Ala	Lys	Asp	Trp 170	Thr	Lys	Gln	Pro	Pro 175	His
Val	Pro	Pro	Leu 180	Tyr	Val	His	Ser	Tyr 185	Thr	Ser	Phe	Ser	Asn 190	Thr	Ile
Ser	Ser	Trp 195	Cys	Gln	Tyr	Glu	Ala 200	Leu	Lys	Tyr	Val	Ser 205	Phe	Pro	Thr
Gln	Thr 210	Ile	Cys	Lys	Ala	Ser 215	Lys	Val	Val	Val	Thr 220	Met	Leu	Met	Gly
Arg 225	Leu	Val	Arg	Gly	Gln 230	Arg	Tyr	Ser	Trp	Phe 235	Glu	Tyr	Gly	Суз	Gly 240
Cys	Thr	Ile	Ala	Phe 245	Gly	Ala	Ser	Leu	Phe 250	Leu	Leu	Ser	Ser	Ser 255	Ser
Lys	Gly	Ala	Gly 260	Ser	Thr	Ile	Thr	Tyr 265	Thr	Ser	Phe	Ser	Gly 270	Met	Ile
Leu	Met	Ala 275	Gly	Tyr	Leu	Leu	Phe 280	Asp	Ala	Phe	Thr	Leu 285	Asn	Trp	Gln
Lys	Ala 290	Leu	Phe	Asp	Thr	Lys 295	Pro	Lys	Val	Ser	Lys 300	Tyr	Gln	Met	Met
Phe 305	Gly	Val	Asn	Phe	Phe 310	Ser	Ala	Ile	Leu	Cys 315	Ala	Val	Ser	Leu	Ile 320
Glu	Gln	Gly	Thr	Leu 325	Trp	Ser	Ser	Ile	Lys 330	Phe	Gly	Ala	Glu	His	Val

Asp Phe Ser Arg Asp Val Phe Leu Leu Ser Leu Ser Gly Ala Ile Gly 340 345 350

Gln Ile Phe Ile Tyr Ser Thr Ile Glu Arg Phe Gly Pro Ile Val Phe 355 360 365

Ala Val Ile Met Thr Ile Arg Gln Ile Phe Ile Arg Asn Thr Leu Ile 370 375 380

Arg Ala Glu Asp His Arg Gly Val Glu Met Ala Pro Pro Pro Pro 385 390 395 400

Glu Pro Phe Arg Leu Lys Phe Leu Ser Met Ile Ile Ala Val Ile His 405 410 415

Ile

<210> 133

<211> 124

<212> PRT

<213> Mus musculus

<400> 133

Met Asp Ala Arg Trp Trp Ala Val Val Leu Ala Thr Leu Pro Ser 1 5 10 15

Leu Gly Ala Gly Glu Ser Pro Glu Ala Pro Pro Gln Ser Trp Thr
20 25 30

Gln Leu Trp Leu Phe Arg Phe Leu Leu Asn Val Ala Gly Tyr Ala Ser 35 40 45

Phe Met Val Pro Gly Tyr Leu Leu Val Gln Tyr Leu Arg Arg Lys Asn 50 55 60

Tyr Leu Glu Thr Gly Arg Gly Leu Cys Phe Pro Leu Val Lys Ala Cys 65 70 75 80

Val Phe Gly Asn Glu Pro Lys Ala Pro Asp Glu Val Leu Leu Ala Pro 85 90 95

Arg Thr Glu Thr Ala Glu Ser Thr Pro Ser Trp Gln Val Leu Lys Leu
100 105 110

Val Phe Cys Ala Ser Gly Leu Gln Thr Gln Phe Leu

<210> 134

<211> 286

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: DUF6 domain sequence

<400> 134

Ser Ser Ala Lys Asn Ala Phe Lys Lys Cys Phe Lys Ser Ile Phe Ser 1 5 10 15

Trp His Asn Glu Thr Val Asn Ile Trp Thr Tyr Lys Lys Glu Lys Phe
20 25 30

Leu Glu Arg Leu Val Lys Leu Ser His Leu Leu Gly Phe Ile Leu Phe 35 40 45

Phe Leu Leu Ile Leu Asp Phe Leu Phe Leu Val Pro Ile Leu Ala 50 55 60

Ser Val Thr Ser His Leu Tyr Ile Leu Gln Asp Arg Val Val Phe Gly 65 70 75 80

Phe Phe Thr Asp Leu Cys Val His Asp Leu Ala Gly Trp Pro Phe Tyr 85 90 95

Phe Leu Gly Ala Phe Leu Cys Leu Leu Leu Ser Ser Ile Tyr His Thr
100 105 110

Phe Ser Cys His Ser Leu Glu Lys Val Ser Glu Phe Phe Leu Lys Leu 115 120 125

Asp Tyr Leu Gly Ile Ser Leu Leu Ile Val Ala Ser Phe Ile Pro Ile 130 135 140

Ile Tyr Tyr Ala Phe Tyr Cys His Pro Phe Phe Arg Thr Leu Tyr Ile 145 150 155 160

Ser Ile Ile Leu Val Leu Gly Leu Ile Ala Ile Tyr Val Ser Leu Ser 165 170 175

Asp Lys Phe Ser Ser Pro Lys Phe Arg Lys Arg Arg Val Pro Leu Arg 180 185 190 Ala Gly Phe Phe Val Leu Leu Gly Leu Ser Gly Val Ile Pro Leu Leu
195 200 205

His Ala Leu Ile Leu Phe Gly Gly His Glu Asn Leu Lys Val Arg Île 210 215 220

Ala Leu Pro Trp Val Leu Leu Met Ala Leu Leu Tyr Ile Val Gly Ala 225 230 235 240

Val Phe Tyr Gly Thr Arg Ile Pro Glu Arg Phe Phe Arg Cys Pro His \$245\$ 250 255

Ala Gly Lys Phe Asp Ile Val Gly His Ser His Gln Leu Phe His Val 260 265 270

Leu Val Val Leu Ala Ala Phe Cys His Tyr Arg Ala Val Leu 275 280 285

<210> 135

<211> 551

<212> PRT

<213> Homo sapiens

<400> 135

Met Leu Pro Leu Leu Leu Pro Leu Leu Trp Gly Gly Ser Leu Gln 1 5 10 15

Glu Lys Pro Val Tyr Glu Leu Gln Val Gln Lys Ser Val Thr Val Gln
20 25 30

Glu Gly Leu Cys Val Leu Val Pro Cys Ser Phe Ser Tyr Pro Trp Arg
35 40 45

Ser Trp Tyr Ser Ser Pro Pro Leu Tyr Val Tyr Trp Phe Arg Asp Gly
50 55 60

Glu Ile Pro Tyr Tyr Ala Glu Val Val Ala Thr Asn Asn Pro Asp Arg
65 70 75 80

Arg Val Lys Pro Glu Thr Gln Gly Arg Phe Arg Leu Leu Gly Asp Val 85 90 95

Gln Lys Lys Asn Cys Ser Leu Ser Ile Gly Asp Ala Arg Met Glu Asp 100 105 110

Thr Gly Ser Tyr Phe Phe Arg Val Glu Arg Gly Arg Asp Val Lys Tyr

		115					120			125				
Ser	Tyr	Gln	Gln	Asn	Lys	Leu	Asn	Leu	Glu	Val	Thr	Ala		
	130					135					140			

Leu Ile Glu

Lys Pro Asp Ile His Phe Leu Glu Pro Leu Glu Ser Gly Arg Pro Thr

Arg Leu Ser Cys Ser Leu Pro Gly Ser Cys Glu Ala Gly Pro Pro Leu

Thr Phe Ser Trp Thr Gly Asn Ala Leu Ser Pro Leu Asp Pro Glu Thr

Thr Arg Ser Ser Glu Leu Thr Leu Thr Pro Arg Pro Glu Asp His Gly

Thr Asn Leu Thr Cys Gln Met Lys Arg Gln Gly Ala Gln Val Thr Thr

Glu Arg Thr Val Gln Leu Asn Val Ser Tyr Ala Pro Gln Thr Ile Thr

Ile Phe Arg Asn Gly Ile Ala Leu Glu Ile Leu Gln Asn Thr Ser Tyr

Leu Pro Val Leu Glu Gly Gln Ala Leu Arg Leu Leu Cys Asp Ala Pro

Ser Asn Pro Pro Ala His Leu Ser Trp Phe Gln Gly Ser Pro Ala Leu

Asn Ala Thr Pro Ile Ser Asn Thr Gly Ile Leu Glu Leu Arg Arg Val

Arg Ser Ala Glu Glu Gly Phe Thr Cys Arg Ala Gln His Pro Leu

Gly Phe Leu Gln Ile Phe Leu Asn Leu Ser Val Tyr Ser Leu Pro Gln

Leu Leu Gly Pro Ser Cys Ser Trp Glu Ala Glu Gly Leu His Cys Arg

Cys Ser Phe Arg Ala Arg Pro Ala Pro Ser Leu Cys Trp Arg Leu Glu

Glu Lys Pro Leu Glu Gly Asn Ser Ser Gln Gly Ser Phe Lys Val Asn

370 375 380

Ser Ser Ser Ala Gly Pro Trp Ala Asn Ser Ser Leu Ile Leu His Gly
385 390 395 400

Gly Leu Ser Ser Asp Leu Lys Val Ser Cys Lys Ala Trp Asn Ile Tyr 405 410 415

Gly Ser Gln Ser Gly Ser Val Leu Leu Gln Gly Arg Ser Asn Leu
420 425 430

Gly Thr Gly Val Val Pro Ala Ala Leu Gly Gly Ala Gly Val Met Ala 435 440 445

Leu Leu Cys Ile Cys Leu Cys Leu Ile Phe Phe Leu Ile Val Lys Ala 450 455 460

Arg Arg Lys Gln Ala Ala Gly Arg Pro Glu Lys Met Asp Asp Glu Asp 465 470 475 480

Pro Ile Met Gly Thr Ile Thr Ser Gly Ser Arg Lys Lys Pro Trp Pro 485 490 495

Asp Ser Pro Gly Asp Gln Ala Ser Pro Pro Gly Asp Ala Pro Pro Leu
500 505 510

Glu Glu Gln Lys Glu Leu His Tyr Ala Ser Leu Ser Phe Ser Glu Met 515 520 525

Lys Ser Arg Glu Pro Lys Asp Gln Glu Ala Pro Ser Thr Thr Glu Tyr 530 540

Ser Glu Ile Lys Thr Ser Lys 545 550

<210> 136

<211> 551

<212> PRT

<213> Homo sapiens

<400> 136

Met Leu Pro Leu Leu Leu Pro Leu Leu Trp Gly Gly Ser Leu Gln
1 5 10 15

Glu Lys Pro Val Tyr Glu Leu Gln Val Gln Lys Ser Val Thr Val Gln 20 25 30

Glu Gly Leu Cys Val Leu Val Pro Cys Ser Phe Ser Tyr Pro Trp Arg Ser Trp Tyr Ser Ser Pro Pro Leu Tyr Val Tyr Trp Phe Arg Asp Gly Glu Ile Pro Tyr Tyr Ala Glu Val Val Ala Thr Asn Asn Pro Asp Arg Arg Val Lys Pro Glu Thr Gln Gly Arg Phe Arg Leu Leu Gly Asp Val Gln Lys Lys Asn Cys Ser Leu Ser Ile Gly Asp Ala Arg Met Glu Asp Thr Gly Ser Tyr Phe Phe Arg Val Glu Arg Gly Arg Asp Val Lys Tyr Ser Tyr Gln Gln Asn Lys Leu Asn Leu Glu Val Thr Ala Leu Ile Glu Lys Pro Asp Ile His Phe Leu Glu Pro Leu Glu Ser Gly Arg Pro Thr Arg Leu Ser Cys Ser Leu Pro Gly Ser Cys Glu Ala Gly Pro Pro Leu Thr Phe Ser Trp Thr Gly Asn Ala Leu Ser Pro Leu Asp Pro Glu Thr Thr Arg Ser Ser Glu Leu Thr Leu Thr Pro Arg Pro Glu Asp His Gly Thr Asn Leu Thr Cys Gln Met Lys Arg Gln Gly Ala Gln Val Thr Thr Glu Arg Thr Val Gln Leu Asn Val Ser Tyr Ala Pro Gln Thr Ile Thr Ile Phe Arg Asn Gly Ile Ala Leu Glu Ile Leu Gln Asn Thr Ser Tyr Leu Pro Val Leu Glu Gly Gln Ala Leu Arg Leu Leu Cys Asp Ala Pro Ser Asn Pro Pro Ala His Leu Ser Trp Phe Gln Gly Ser Pro Ala Leu

Asn Ala Thr Pro Ile Ser Asn Thr Gly Ile Leu Glu Leu Arg Arg Val Arg Ser Ala Glu Lys Gly Gly Phe Thr Cys Arg Ala Gln His Pro Leu Gly Phe Leu Gln Ile Phe Leu Asn Leu Ser Val Tyr Ser Leu Pro Gln Leu Leu Gly Pro Ser Cys Ser Trp Glu Ala Glu Gly Leu His Cys Arg Cys Ser Phe Arg Ala Trp Pro Ala Pro Ser Leu Cys Trp Arg Leu Glu Glu Lys Pro Leu Glu Gly Asn Ser Ser Gln Gly Ser Phe Lys Val Asn Ser Ser Ser Pro Gly Pro Trp Ala Asn Ser Ser Leu Ile Leu His Gly Gly Leu Asn Ser Asp Leu Lys Val Ser Cys Lys Ala Trp Asn Ile Tyr Gly Ser Gln Ser Gly Ser Val Leu Leu Gln Gly Arg Ser Asn Leu Gly Thr Gly Val Val Pro Ala Ala Leu Gly Gly Ala Gly Val Met Ala Leu Leu Cys Ile Cys Leu Cys Leu Ile Phe Phe Leu Ile Val Lys Ala Arg Arg Lys Gln Ala Ala Gly Arg Pro Glu Lys Met Asp Asp Glu Asp Pro Ile Met Gly Thr Ile Thr Ser Gly Ser Arg Lys Lys Pro Trp Pro Asp Ser Pro Gly Asp Gln Ala Ser Pro Pro Gly Asp Ala Pro Pro Leu Glu Glu Gln Lys Glu Leu His Tyr Ala Ser Leu Ser Phe Ser Glu Met Lys Ser Arg Glu Pro Lys Asp Gln Glu Ala Pro Ser Thr Thr Glu Tyr

Ser Glu Ile Lys Thr Ser Lys <210> 137 <211> 442 <212> PRT <213> Homo sapiens <400> 137 Met Leu Pro Leu Leu Pro Leu Leu Trp Ala Gly Ala Leu Ala Gln Glu Arg Arg Phe Gln Leu Glu Gly Pro Glu Ser Leu Thr Val Gln Glu Gly Leu Cys Val Leu Val Pro Cys Arg Leu Pro Thr Thr Leu Pro Ala Ser Tyr Tyr Gly Tyr Gly Tyr Trp Phe Leu Glu Gly Ala Asp Val Pro Val Ala Thr Asn Asp Pro Asp Glu Glu Val Gln Glu Glu Thr Arg Gly Arg Phe His Leu Leu Trp Asp Pro Arg Arg Lys Asn Cys Ser Leu Ser Ile Arg Asp Ala Arg Arg Asp Asn Ala Ala Tyr Phe Phe Arg Leu Lys Ser Lys Trp Met Lys Tyr Gly Tyr Thr Ser Ser Lys Leu Ser Val Arg Val Met Ala Leu Thr His Arg Pro Asn Ile Ser Ile Pro Gly Thr Leu Glu Ser Gly His Pro Ser Asn Leu Thr Cys Ser Val Pro Trp Val

Thr Ser Leu Gly Pro Arg Thr Thr Gln Ser Ser Val Leu Thr Ile Thr Pro Arg Pro Gln Asp His Ser Thr Asn Leu Thr Cys Gln Val Thr Phe

Cys Glu Gln Gly Thr Pro Pro Ile Phe Ser Trp Met Ser Ala Ala Pro

Pro Gly Ala Gly Val Thr Met Glu Arg Thr Ile Gln Leu Asn Val Ser Tyr Ala Pro Gln Lys Val Ala Ile Ser Ile Phe Gln Gly Asn Ser Ala Ala Phe Lys Ile Leu Gln Asn Thr Ser Ser Leu Pro Val Leu Glu Gly Gln Ala Leu Arg Leu Leu Cys Asp Ala Asp Gly Asn Pro Pro Ala His Leu Ser Trp Phe Gln Gly Phe Pro Ala Leu Asn Ala Thr Pro Ile Ser Asn Thr Gly Val Leu Glu Leu Pro Gln Val Gly Ser Ala Glu Glu Gly Asp Phe Thr Cys Arg Ala Gln His Pro Leu Gly Ser Leu Gln Ile Ser Leu Ser Leu Phe Val His Trp Lys Pro Glu Gly Arg Ala Gly Gly Val Leu Gly Ala Val Trp Gly Ala Ser Ile Thr Thr Leu Val Phe Leu Cys Val Cys Phe Ile Phe Arg Val Lys Thr Arg Arg Lys Lys Ala Ala Gln Pro Val Gln Asn Thr Asp Asp Val Asn Pro Val Met Val Ser Gly Ser Arg Gly His Gln His Gln Phe Gln Thr Gly Ile Val Ser Asp His Pro Ala Glu Ala Gly Pro Ile Ser Glu Asp Glu Gln Glu Leu His Tyr Ala Val Leu His Phe His Lys Val Gln Pro Gln Glu Pro Lys Val Thr Asp Thr Glu Tyr Ser Glu Ile Lys Ile His Lys

<211> 440

<212> PRT

<213> Homo sapiens

<400> 138

Met Leu Pro Leu Leu Pro Leu Leu Trp Ala Gly Ala Leu Ala Gln
1 5 10 15

Glu Arg Arg Phe Gln Leu Glu Gly Pro Glu Ser Leu Thr Val Gln Glu 20 25 30

Gly Leu Cys Val Leu Val Pro Cys Arg Leu Pro Thr Thr Leu Pro Ala 35 40 45

Ser Tyr Tyr Gly Tyr Gly Tyr Trp Phe Leu Glu Gly Ala Asp Val Pro 50 55 60

Val Ala Thr Asn Asp Pro Asp Glu Glu Val Gln Glu Glu Thr Arg Gly
65 70 75 80

Arg Phe His Leu Leu Trp Asp Pro Arg Arg Lys Asn Cys Ser Leu Ser 85 90 95

Ile Arg Asp Ala Arg Arg Arg Asp Asn Ala Ala Tyr Phe Phe Arg Leu
100 105 110

Lys Ser Lys Trp Met Lys Tyr Gly Tyr Thr Ser Ser Lys Ile Tyr Val 115 120 125

Arg Val Met Ala Leu Thr His Arg Pro Asn Ile Ser Ile Pro Gly Pro 130 135 140

Gly Val Trp Pro Ser Ser Asn Leu Thr Cys Ser Val Pro Trp Val Cys 145 150 155 160

Glu Gln Gly Thr Pro Pro Ile Phe Ser Trp Met Ser Ala Ala Pro His 165 170 175

Leu Leu Gly Pro Arg Thr Thr Gln Ser Ser Val Leu Thr Ile Thr Pro
180 185 190

Ala Gln Asp His Ser Thr Asn Leu Thr Cys Gln Val Thr Phe Pro Gly
195 200 205

Ala Gly Val Thr Met Glu Arg Thr Ile Gln Leu Asn Val Ser Tyr Ala 210 215 220

Pro Gln Lys Val Ala Ile Ser Ile Phe Gln Gly Asn Ser Ala Ala Phe

Lys Ile Leu Gln Asn Thr Ser Ser Leu Pro Val Leu Glu Gly Gln Ala 245 250 255

Leu Arg Leu Cys Asp Ala Asp Gly Asn Pro Pro Ala His Leu Ser 260 265 270

Trp Phe Gln Gly Phe Pro Ala Leu Asn Ala Thr Pro Ile Ser Asn Thr 275 280 285

Gly Val Leu Glu Leu Pro Gln Val Gly Ser Ala Glu Glu Gly Asp Phe 290 295 300

Thr Cys Arg Ala Gln His Pro Leu Gly Ser Leu Gln Ile Ser Leu Ser 305 310 315 320

Leu Phe Val His Trp Lys Pro Glu Gly Arg Ala Gly Gly Val Leu Gly 325 330 335

Ala Val Trp Gly Ala Ser Ile Thr Thr Leu Val Phe Leu Cys Val Cys 340 345 350

Phe Ile Phe Arg Val Lys Thr Arg Arg Lys Lys Ala Ala Gln Pro Val 355 360 365

Gln Asn Thr Asp Asp Val Asn Pro Val Met Val Ser Gly Ser Arg Gly 370 375 380

His Gln His Gln Phe Gln Thr Gly Ile Val Ser Asp His Pro Ala Glu 385 390 395 400

Ala Gly Pro Ile Ser Glu Asp Glu Gln Glu Leu His Tyr Ala Val Leu 405 410 415

His Phe His Lys Val Gln Pro Gln Glu Pro Lys Val Thr Asp Thr Glu 420 425 430

Tyr Ser Glu Ile Lys Ile His Lys 435 440

<210> 139

<211> 463

<212> PRT

<213> Homo sapiens

<400> 139

Met 1	Leu	Leu	Leu	Leu 5	Leu	Pro	Leu	Leu	Trp 10	Gly	Arg	Glu	Arg	Ala 15	Glu
Gly	Gln	Thr	Ser 20	Lys	Leu	Leu	Thr	Met 25	Gln	Ser	Ser	Val	Thr 30	Val	Gln
Glu	Gly	Leu 35	Cys	Val	His	Val	Pro 40	Cys	Ser	Phe	Ser	Tyr 45	Pro	Ser	His
Gly	Trp 50	Ile	Tyr	Pro	Gly	Pro 55	Val	Val	His	Gly	Tyr 60	Trp	Phe	Arg	Glu
Gly 65	Ala	Asn	Thr	Asp	Gln 70	Asp	Ala	Pro	Val	Ala 75	Thr	Asn	Asn	Pro	Ala 80
Arg	Ala	Val	Trp	Glu 85	Glu	Thr	Arg	Asp	Arg 90	Phe	His	Leu	Leu	Gly 95	Asp
Pro	His	Thr	Glu 100	Asn	Cys	Thr	Leu	Ser 105	Ile	Arg	Asp	Ala	Arg 110	Arg	Ser
Asp	Ala	Gly 115	Arg	Tyr	Phe	Phe	Arg 120	Met	Glu	Lys	Gly	Ser 125	Ile	Lys	Trp
Asn	Tyr 130	Lys	His	His	Arg	Leu 135	Ser	Val	Asn	Val	Thr 140	Ala	Leu	Thr	His
Arg 145	Pro	Asn	Ile	Leu	Ile 150	Pro	Gly	Thr	Leu	Glu 155	Ser	Gly	Cys	Pro	Gln 160
Asn	Leu	Thr	Cys	Ser 165	Val	Pro	Trp	Ala	Cys 170	Glu	Gln	Gly	Thr	Pro 175	Pro
Met	Ile	Ser	Trp 180	Ile	Gly	Thr	Ser	Val 185	Ser	Pro	Leu	Asp	Pro 190	Ser	Thr
Thr	Arg	Ser 195	Ser	Val	Leu	Thr	Leu 200	Ile	Pro	Gln	Pro	Gln 205	Asp	His	Gly
Thr	Ser 210	Leu	Thr	Cys	Gln	Val 215	Thr	Phe	Pro	Gly	Ala 220	Ser	Val	Thr	Thr
Asn 225	Lys	Thr	Val	His	Leu 230	Asn	Val	Ser	Tyr	Pro 235	Pro	Gln	Asn	Leu	Thr 240
Met	Thr	Val	Phe	Gln 245	Gly	Asp	Gly	Thr	Val 250	Ser	Thr	Val	Leu	Gly 255	Asn

Gly Ser Ser Leu Ser Leu Pro Glu Gly Gln Ser Leu Arg Leu Val Cys 260 265 270

Ala Val Asp Ala Val Asp Ser Asn Pro Pro Ala Arg Leu Ser Leu Ser 275 280 285

Trp Arg Gly Leu Thr Leu Cys Pro Ser Gln Pro Ser Asn Pro Gly Val 290 295 300

Leu Glu Leu Pro Trp Val His Leu Arg Asp Glu Ala Glu Phe Thr Cys 305 310 315 320

Arg Ala Gln Asn Pro Leu Gly Ser Gln Gln Val Tyr Leu Asn Val Ser 325 330 335

Leu Gln Ser Lys Ala Thr Ser Gly Val Thr Gln Gly Val Val Gly Gly 340 345 350

Ala Gly Ala Thr Ala Leu Val Phe Leu Ser Phe Cys Val Ile Phe Val 355 360 365

Val Val Arg Ser Cys Arg Lys Lys Ser Ala Arg Pro Ala Ala Gly Val 370 375 380

Gly Asp Thr Gly Ile Glu Asp Ala Asn Ala Val Arg Gly Ser Ala Ser 385 390 395 400

Gln Gly Pro Leu Thr Glu Pro Trp Ala Glu Asp Ser Pro Pro Asp Gln
405 410 415

Pro Pro Pro Ala Ser Ala Arg Ser Ser Val Gly Glu Gly Glu Leu Gln 420 425 430

Tyr Ala Ser Leu Ser Phe Gln Met Val Lys Pro Trp Asp Ser Arg Gly 435 440 445

Gln Glu Ala Thr Asp Thr Glu Tyr Ser Glu Ile Lys Ile His Arg 450 455 460

<210> 140

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Immunoglobin domain sequence

<400> 140

Ser Val Ser Gly Phe Gly Pro Pro Pro Val Thr Trp Leu Arg Asn Gly
1 5 10 15

Lys Leu Ser Leu Thr Ile Ser Val Thr Pro Glu Asp Ser Gly Gly Thr 20 25 30

Tyr Thr

<210> 141

<211> 290

<212> PRT

<213> Homo sapiens

<400> 141

Ala His Thr Glu Tyr Pro Val Asn Thr Ile Ile Ile His Glu Asp Phe 1 5 10 15

Asp Asn Asn Ser Met Ser Asn Asn Ile Ala Leu Leu Lys Thr Asp Thr 20 25 30

Ala Met His Phe Gly Asn Leu Val Gln Ser Ile Cys Phe Leu Gly Arg 35 40 45

Met Leu His Thr Pro Pro Val Leu Gln Asn Cys Trp Val Ser Gly Trp 50 55 60

Asn Pro Thr Ser Ala Thr Gly Asn His Met Thr Met Ser Val Leu Arg
65 70 75 80

Lys Ile Phe Val Lys Asp Leu Asp Met Cys Pro Leu Tyr Lys Leu Gln 85 90 95

Lys Thr Glu Cys Gly Ser His Thr Lys Glu Glu Thr Lys Thr Ala Cys 100 105 110

Leu Gly Asp Pro Gly Ser Pro Met Met Cys Gln Leu Gln Gln Phe Asp · 115 120 125

Leu Trp Val Leu Arg Gly Ile Leu Asn Phe Gly Gly Glu Thr Cys Pro 130 135 140

Gly Leu Phe Leu Tyr Thr Lys Val Glu Asp Tyr Ser Lys Trp Ile Thr 145 150 155 160

Ser Lys Ala Glu Arg Ala Gly Pro Pro Leu Ser Ser Leu His His Trp Glu Lys Leu Ile Ser Phe Ser His His Gly Pro Asn Ala Ala Met Thr Gln Lys Thr Tyr Ser Asp Ser Glu Leu Gly His Val Gly Ser Tyr Leu Gln Gly Gln Arg Arg Thr Ile Thr His Ser Arg Leu Gly Asn Ser Ser Arg Asp Ser Leu Asp Val Arg Glu Lys Asp Val Lys Glu Ser Gly Arg Ser Pro Glu Ala Ser Val Gln Pro Leu Tyr Tyr Asp Tyr Tyr Gly Gly Glu Val Gly Glu Gly Arg Ile Phe Ala Gly Gln Asn Arg Leu Tyr Gln Pro Glu Glu Ile Ile Leu Val Ser Phe Val Leu Val Phe Phe Cys Ser Ser Ile <210> 142 <211> 270 <212> PRT <213> Homo sapiens <400> 142 Met Ser Asn Asn Ile Ala Leu Leu Lys Thr Asp Thr Ala Met His Phe Gly Asn Leu Val Gln Ser Ile Cys Phe Leu Gly Arg Met Leu His Thr Pro Pro Val Leu Gln Asn Cys Trp Val Ser Gly Trp Asn Pro Thr Ser Ala Thr Gly Asn His Met Thr Met Ser Val Leu Arg Lys Ile Phe Val Lys Asp Leu Asp Met Cys Pro Leu Tyr Lys Leu Gln Lys Thr Glu Cys

Gly Ser His Thr Lys Glu Glu Thr Lys Thr Ala Cys Leu Gly Asp Pro 85 90 95

Gly Ser Pro Met Met Cys Gln Leu Gln Gln Phe Asp Leu Trp Val Leu

100 105 110

Arg Gly Val Leu Asn Phe Gly Gly Glu Thr Cys Pro Gly Leu Phe Leu 115 120 125

Tyr Thr Lys Val Glu Asp Tyr Ser Lys Trp Ile Thr Ser Lys Ala Glu 130 140

Arg Ala Gly Pro Pro Leu Ser Ser Leu His His Trp Glu Lys Leu Ile 145 150 155 160

Ser Phe Ser His His Gly Pro Asn Ala Thr Met Thr Gln Lys Thr Tyr 165 170 175

Ser Asp Ser Glu Leu Gly His Val Gly Ser Tyr Leu Gln Gly Gln Arg \$180\$ \$185\$ \$190\$

Arg Thr Ile Thr His Ser Arg Leu Gly Asn Ser Ser Arg Asp Ser Leu 195 200 205

Asp Val Arg Glu Lys Asp Val Lys Glu Ser Gly Arg Ser Pro Glu Ala 210 215 220

Ser Val Gln Pro Leu Tyr Tyr Asp Tyr Tyr Gly Gly Glu Val Gly Glu 225 230 235 240

Gly Arg Ile Phe Ala Gly Gln Asn Arg Leu Tyr Gln Pro Glu Glu Ile 245 250 255

Ile Leu Val Ser Phe Val Leu Val Phe Phe Cys Ser Ser Ile 260 265 270

<210> 143

<211> 624

<212> PRT

<213> Mus musculus

<400> 143

Met Thr Ser Leu His Gln Val Leu Tyr Phe Ile Phe Phe Ala Ser Val 1 5 10 15

Ser Ser Glu Cys Val Thr Lys Val Phe Lys Asp Ile Ser Phe Gln Gly

20

25

- Gly Asp Leu Ser Thr Val Phe Thr Pro Ser Ala Thr Tyr Cys Arg Leu
 35 40 45
- Val Cys Thr His His Pro Arg Cys Leu Leu Phe Thr Phe Met Ala Glu 50 55 60
- Ser Ser Ser Asp Asp Pro Thr Lys Trp Phe Ala Cys Ile Leu Lys Asp 65 70 75 80
- Ser Val Thr Glu Ile Leu Pro Met Val Asn Met Thr Gly Ala Ile Ser 85 90 95
- Gly Tyr Ser Phe Lys Gln Cys Pro Gln Gln Leu Ser Thr Cys Ser Lys 100 105 110
- Asp Val Tyr Val Asn Leu Asp Met Lys Gly Met Asn Tyr Asn Ser Ser 115 120 125
- Val Val Lys Asn Ala Arg Glu Cys Gln Glu Arg Cys Thr Asp Asp Ala 130 135 140
- His Cys Gln Phe Phe Thr Tyr Ala Thr Gly Tyr Phe Pro Ser Val Asp 145 150 155 160
- His Arg Lys Met Cys Leu Leu Lys Tyr Thr Arg Thr Gly Thr Pro Thr 165 170 175
- Thr Ile Thr Lys Leu Asn Gly Val Val Ser Gly Phe Ser Leu Lys Ser 180 185 190
- Cys Gly Leu Ser Asn Leu Ala Cys Ile Arg Asp Ile Phe Pro Asn Thr 195 200 205
- Val Leu Ala Asp Leu Asn Ile Asp Ser Val Val Ala Pro Asp Ala Phe 210 215 220
- Val Cys Arg Arg Ile Cys Thr His His Pro Thr Cys Leu Phe Phe Thr 225 230 235 240
- Phe Phe Ser Gln Ala Trp Pro Lys Glu Ser Gln Arg His Leu Cys Leu 245 250 255
- Leu Lys Thr Ser Glu Ser Gly Leu Pro Ser Thr Arg Ile Thr Lys Ser 260 265 270
- His Ala Leu Ser Gly Phe Ser Leu Gln His Cys Arg His Ser Val Pro

Val	Phe 290	Cys	His	Pro	Ser	Phe 295	Tyr	Asn	Asp	Thr	Asp 300	Phe	Leu	Gly	Glu
Glu 305	Leu	Asp	Ile	Val	Asp 310	Val	Lys	Gly	Gln	Glu 315	Thr	Cys	Gln	Lys	Thr 320
Cys	Thr	Asn	Asn	Ala 325	Arg	Cys	Gln	Phe	Phe 330	Thr	Tyr	Tyr	Pro	Ser 335	His
Arg	Leu	Cys	Asn 340	Glu	Arg	Asn	Arg	Arg 345	Gly	Arg	Cys	Tyr	Leu 350	Lys	Leu
Ser	Ser	Asn 355	Gly	Ser	Pro	Thr	Arg 360	Ile	Leu	His	Gly	Arg 365	Gly	Gly	Ile
Ser	Gly 370	Tyr	Ser	Leu	Arg	Leu 375	Cys	Lys	Met	Asp	Asn 380	Val	Cys	Thr	Thr
Lys 385	Ile	Asn	Pro	Arg	Val 390	Val	Gly	Gly	Ala	Ala 395	Ser	Val	His	Gly	Glu 400
Trp	Pro	Trp	Gln	Val 405	Thr	Leu	His	Ile	Ser 410	Gln	Gly	His	Leu	Cys 415	Gly
Gly	Ser	Ile	Ile 420	Gly	Asn	Gln	Trp	Ile 425	Leú	Thr	Ala	Ala	His 430	Cys	Phe
Ser	Gly	Ile 435	Glu	Thr	Pro	Lys	Lys 440	Leu	Arg	Val	Tyr	Gly 445	Gly	Ile	Val
Asn	Gln 450	Ser	Glu	Ile	Asn	Glu 455	Gly	Thr	Ala	Phe	Phe 460	Arg	Val	Gln	Glu
Met 465	Ile	Ile	His	Asp	Gln 470	Tyr	Thr	Thr	Ala	Glu 475	Ser	Gly	Tyr	Asp	Ile 480
Ala	Leu	Leu	Lys	Leu 485	Glu	Ser	Ala	Met	Asn 490	Tyr	Thr	Asp	Phe	Gln 495	Arg
Pro	Ile	Cys	Leu 500	Pro	Ser	Lys	Gly	Asp 505	Arg	Asn	Ala	Val	His 510	Thr	Glu
Cys	Trp	Val	Thr	Gly	Trp	Gly	Tyr 520	Thr	Ala	Leu	Arg	Gly	Glu	Val	Gln

Ser Thr Leu Gln Lys Ala Lys Val Pro Leu Val Ser Asn Glu Glu Cys

530 535 540

Gln Thr Arg Tyr Arg Arg His Lys Ile Thr Asn Lys Met Ile Cys Ala 545 550 555 560

Gly Tyr Lys Glu Gly Gly Lys Asp Thr Cys Lys Gly Asp Ser Gly Gly 565 570 575

Pro Leu Ser Cys Lys Tyr Asn Gly Val Trp His Leu Val Gly Ile Thr 580 585 590

Ser Trp Gly Glu Gly Cys Gly Gln Lys Glu Arg Pro Gly Val Tyr Thr 595 600 605

Asn Val Ala Lys Tyr Val Asp Trp Ile Leu Glu Lys Thr Gln Thr Val 610 620

<210> 144

<211> 326

<212> PRT

<213> Mus musculus

<400> 144

Met Cys Arg Gln Pro Met Lys Arg Trp Lys Asp Arg Arg Thr Gly Leu 1 5 10 15

Leu Leu Pro Leu Val Leu Leu Leu Phe Gly Ala Cys Ser Ser Leu Ala 20 25 30

Trp Val Cys Gly Arg Arg Met Ser Ser Arg Ser Gln Gln Leu Asn Asn 35 40 45

Ala Ser Ala Ile Val Glu Gly Lys Pro Ala Ser Ala Ile Val Gly Gly 50 55 60

Lys Pro Ala Asn Ile Leu Glu Phe Pro Trp His Val Gly Ile Met Asn 65 70 75 80

His Gly Ser His Leu Cys Gly Gly Ser Ile Leu Asn Glu Trp Trp Val 85 90 95

Leu Ser Ala Ser His Cys Phe Asp Gln Leu Asn Asn Ser Lys Leu Glu 100 105 110 Ile Ile His Gly Thr Glu Asp Leu Ser Thr Lys Gly Ile Lys Tyr Gln
115 120 125

Lys Val Asp Lys Leu Phe Leu His Pro Lys Phe Asp Asp Trp Leu Leu 130 135 140

Asp Asn Asp Ile Ala Leu Leu Leu Leu Lys Ser Pro Leu Asn Leu Ser 145 150 155 160

Val Asn Arg Ile Pro Ile Cys Thr Ser Glu Ile Ser Asp Ile Gln Ala 165 170 175

Trp Arg Asn Cys Trp Val Thr Gly Trp Gly Ile Thr Asn Thr Ser Glu
180 185 190

Lys Gly Val Gln Pro Thr Ile Leu Gln Ala Val Lys Val Asp Leu Tyr 195 200 205

Arg Trp Asp Trp Cys Gly Tyr Ile Leu Ser Leu Leu Thr Lys Asn Met 210 215 220

Leu Cys Ala Gly Thr Gln Asp Pro Gly Lys Asp Ala Cys Gln Gly Asp 225 230 235 240

Ser Gly Gly Ala Leu Val Cys Asn Lys Lys Arg Asn Thr Ala Ile Trp 245 250 255

Tyr Gln Val Gly Ile Val Ser Trp Gly Met Gly Cys Gly Lys Lys Asn 260 265 270

Leu Pro Gly Val Tyr Thr Lys Val Ser His Tyr Val Arg Trp Ile Ser 275 280 285

Lys Gln Thr Ala Lys Ala Gly Arg Pro Tyr Met Tyr Glu Gln Asn Ser 290 295 300

Ala Cys Pro Leu Val Leu Ser Cys Arg Ala Ile Leu Phe Leu Tyr Phe 305 310 315 320

Val Met Phe Leu Leu Thr 325

<210> 145

<211> 624

<212> PRT

<213> Mus musculus

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Ser	Ser	Glu	Cys 20	Val	Thr	Lys	Val	Phe 25	Lys	Asp	Ile	Ser	Phe 30	Gln	Gly
Gly	Asp	Leu 35	Ser	Thr	Val	Phe	Thr 40	Pro	Ser	Ala	Thr	Tyr 45	Cys	Arg	Leu
Val	Cys 50	Thr	His	His	Pro	Arg 55	Cys	Leu	Leu	Phe	Thr 60	Phe	Met	Ala	Glu
Ser 65	Ser	Ser	Asp	Asp	Pro 70	Thr	Lys	Trp	Phe	Ala 75	Cys	Ile	Leu	Lys	Asp 80
Ser	Val	Thr	Glu	Ile 85	Leu	Pro	Met	Val	Asn 90	Met	Thr	Gly	Ala	Ile 95	Ser
Gly	Tyr	Ser	Phe 100	Lys	Gln	Cys	Pro	Gln 105	Gln	Leu	Ser	Thr	Cys 110	Ser	Lys
Asp	Glu	Tyr	Val	Asn	Leu	Asp	Met	Lys	Gly	Met	Asn	Tyr	Asn	Ser	Ser

- Asp Glu Tyr Val Asn Leu Asp Met Lys Gly Met Asn Tyr Asn Ser Ser 115 120 125
- Val Val Lys Asn Ala Arg Glu Cys Gln Glu Arg Cys Thr Asp Asp Ala 130 135 140
- His Arg Lys Met Cys Leu Leu Lys Tyr Thr Arg Thr Gly Thr Pro Thr 165 170 175
- Thr Ile Thr Lys Leu Asn Gly Val Val Ser Gly Phe Ser Leu Lys Ser 180 185 190
- Cys Gly Leu Ser Asn Leu Ala Cys Ile Arg Asp Ile Phe Pro Asn Thr 195 200 205
- Val Leu Ala Asp Leu Asn Ile Asp Ser Val Val Ala Pro Asp Ala Phe 210 215 220
- Val Cys Arg Arg Ile Cys Thr His His Pro Thr Cys Leu Phe Phe Thr 225 230 235 240
- Phe Phe Ser Gln Ala Trp Pro Lys Glu Ser Gln Arg His Leu Cys Leu 245 250 255

Leu Lys Thr Ser Glu Ser Gly Leu Pro Ser Thr Arg Ile Thr Lys Ile His Ala Leu Ser Gly Phe Ser Leu Gln His Cys Arg His Ser Val Pro Val Phe Cys His Pro Ser Phe Tyr Asn Asp Thr Asp Phe Leu Gly Glu Glu Leu Asp Ile Val Asp Val Lys Gly Gln Glu Thr Cys Gln Lys Thr Cys Thr Asn Asn Ala Arg Cys Gln Phe Phe Thr Tyr Tyr Pro Ser His Arg Leu Cys Asn Glu Arg Asn Arg Gly Arg Cys Tyr Leu Lys Leu Ser Ser Asn Gly Ser Pro Thr Arg Ile Leu His Gly Arg Gly Gly Leu Ser Gly Tyr Ser Leu Arg Leu Cys Lys Met Asp Asn Val Cys Thr Thr Lys Ile Asn Pro Arg Val Val Gly Gly Ala Ala Ser Val His Gly Glu Trp Pro Trp Gln Val Thr Leu His Ile Ser Gln Gly His Leu Cys Gly Gly Ser Ile Ile Gly Asn Gln Trp Ile Leu Thr Ala Ala His Cys Phe Ser Gly Ile Glu Thr Pro Lys Lys Leu Arg Val Tyr Gly Gly Ile Val Asn Gln Ser Glu Ile Asn Glu Gly Thr Ala Phe Phe Arg Glu Gln Glu Met Ile Ile His Asp Gln Tyr Thr Thr Ala Glu Ser Gly Tyr Asp Ile Ala Leu Leu Lys Leu Glu Ser Ala Met Asn Tyr Thr Asp Phe Gln Arg Pro Ile Cys Leu Pro Ser Lys Gly Asp Arg Asn Ala Val His Thr Glu

Cys Trp Val Thr Gly Trp Gly Tyr Thr Ala Leu Arg Gly Glu Val Gln 515 520 525

Ser Thr Leu Gln Lys Ala Lys Val Pro Leu Val Ser Asn Glu Glu Cys 530 540

Gln Thr Arg Tyr Arg Arg His Lys Ile Thr Asn Lys Met Ile Cys Ala 545 550 555 560

Gly Tyr Lys Glu Gly Gly Lys Asp Thr Cys Lys Gly Asp Ser Gly Gly 565 570 575

Pro Leu Ser Cys Lys Tyr Asn Gly Val Trp His Leu Val Gly Ile Thr 580 585 590

Ser Trp Gly Glu Gly Cys Gly Gln Lys Glu Arg Pro Gly Val Tyr Thr 595 600 605

Asn Val Ala Lys Tyr Val Asp Trp Ile Leu Glu Lys Thr Gln Thr Val 610 620

<210> 146

<211> 213

<212> PRT

<213> Homo sapiens

<400> 146

Glu Phe Pro Trp Val Val Ser Leu Gln Asp Ser Gln Tyr Thr His Leu

1 5 10 15

Ala Phe Gly Cys Ile Leu Ser Glu Phe Trp Val Leu Ser Ile Ala Ser 20 25 30

Ala Ile Gln Asn Arg Lys Asp Ile Val Val Ile Val Gly Ile Ser Asn 35 40 45

Met Asp Pro Ser Lys Ile Ala His Thr Glu Tyr Pro Val Asn Thr Ile 50 55 60

Ile Ile His Glu Asp Phe Asp Asn Asn Ser Met Ser Asn Asn Ile Ala
65 70 75 80

Leu Leu Lys Thr Asp Thr Ala Met His Phe Gly Asn Leu Val Gln Ser

95

Ile Cys Phe Leu Gly Arg Met Leu His Thr Pro Pro Val Leu Gln Asn
100 105 110

Cys Trp Val Ser Gly Trp Asn Pro Thr Ser Ala Thr Gly Asn His Met 115 120 125

Thr Met Ser Val Leu Arg Lys Ile Phe Val Lys Asp Leu Asp Met Cys 130 135 140

Pro Leu Tyr Lys Leu Gln Lys Thr Glu Cys Gly Ser His Thr Lys Glu 145 150 155 160

Glu Thr Lys Thr Ala Cys Leu Gly Asp Pro Gly Ser Pro Met Met Cys 165 170 175

Gln Leu Gln Gln Phe Asp Leu Trp Val Leu Arg Gly Ile Leu Asn Phe 180 185 190

Gly Gly Glu Thr Cys Pro Gly Leu Phe Leu Tyr Thr Lys Val Glu Asp 195 200 205

Tyr Ser Lys Trp Ile 210

<210> 147

<211> 207

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Trypsin domain sequence

<400> 147

Ser Phe Pro Trp Gln Val Ser Leu Gln Val Ser Ser Gly His Phe Cys
1 10 15

Gly Gly Ser Leu Ile Ser Glu Asn Trp Val Leu Thr Ala Ala His Cys 20 25 30

Val Ser Gly Ala Ser Ser Val Arg Val Val Leu Gly Glu His Asn Leu 35 40 45

Gly Thr Thr Glu Gly Thr Glu Gln Lys Phe Asp Val Lys Lys Ile Ile 50 55 60

Val His Pro Asn Tyr Asn Pro Asp Thr Asn Asp Ile Ala Leu Leu Lys
65 70 75 80

Leu Lys Ser Pro Val Thr Leu Gly Asp Thr Val Arg Pro Ile Cys Leu 85 90 95

Pro Ser Ala Ser Ser Asp Leu Pro Val Gly Thr Thr Cys Ser Val Ser 100 105 110

Gly Trp Gly Arg Thr Lys Asn Leu Gly Thr Ser Asp Thr Leu Gln Glu
115 120 125

Val Val Val Pro Ile Val Ser Arg Glu Thr Cys Arg Ser Ala Tyr Gly
130 135 140

Gly Thr Val Thr Asp Thr Met Ile Cys Ala Gly Ala Leu Gly Gly Lys
145 150 155 160

Asp Ala Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Ser Asp Gly 165 170 175

Glu Leu Val Gly Ile Val Ser Trp Gly Tyr Gly Cys Ala Val Gly Asn 180 185 190

Tyr Pro Gly Val Tyr Thr Arg Val Ser Arg Tyr Leu Asp Trp Ile 195 200 205

<210> 148

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 148

gatccttgga aacaaccaga tc

22

<210> 149

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223>	Description of Artificial sequence	Sequence:	PCR Primer	
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<210>	150			
<211>				
<212>	DNA Artificial Sequence			
\213/	Artificial Sequence			
<220>				
<223>	Description of Artificial	Sequence:	PCR Primer	
	sequence			
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<210>	151			
<211>				
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<213>	Artificial Sequence			
<220>				
	Description of Artificial	Sequence:	PCR Primer	
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	agtgc aacacagata tc			22
			,	
<210>	150			
<211>				
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<220>				
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	4.00			
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ccaugi	gtcta cacaaaagct ttccca			20
<210>				
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<212>	DNA			
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	_			
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12207	sequence	bequence.	TOR TITMET	
	sequence			
	153			
<400>				
gcttc	ctgaa ggttttgttg a			21
<210>	154			
<211>	22			
<212>	DNA			
<213>	Artificial Sequence			
<220>				
	Description of Artificial	Soguengo:	DCD Drimor	
\ 2237		sequence:	PCK PIIMei	
	sequence			
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ggttt	gtgct gcttctaaca tc			22
<210>	155			
<211>	24			
<212>	DNA			
<213>	Artificial Sequence			
	1			
<220>				
	Description of Artificial	Campanaa	DCD Designation	
\223/		sequence:	PCR Primer	
	sequence			
<400>				
acacca	agogg tgotootott caat			24
<210>	156			
<211>	22			
<212>	DNA			
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<220>				
	Description of Table 1.1	Q	DOD D.	
~ ∠∠3>	Description of Artificial	sequence:	rck rrimer	
	sequence			
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cattga	igcat cttacggttt gt			22

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<211> 20
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<223> Description of Artificial Sequence: PCR Primer
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ctgggcatcc agaagatctt
                                                                    20
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<400> 158
ctctgcaagt acagcggcta cctgg
                                                                    25
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cctcgtcatt cagttccagt ac
                                                                    22
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sequence

<400> 160 ggtgccaata cgaagctctt a	21
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<210> 163 <211> 22 <212> DNA <213> Artificial Sequence	
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~Z13>	Artificial Sequence			
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ccagto	caata ccatcatcat ccatgagg			28

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tatgttgttg ctcatggagt tg
                                                                    22
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<400> 169
tggcttattc agaagagcat aaagg
                                                                    25
<210> 170
<211> 27
<212> DNA
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<210> 172 <211> 27		
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agtgactaga gatcctccag gtcagtt		27
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sequence	PCR Primer	
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<400> 174		
cgtcatcctg agcccgtccg tc		22
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٤	sequence			
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gtttcgc	ggcc ctgtgcgg			18
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	Description of Artificial	Sequence:	PCR Primer	
S	sequence			
<400> 1	7.6			
				27
grägräc	cca tttgttttcc tcagagt			21
<210> 1	77			
<211> 2				
<212> D				
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s	equence	_		
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ggtcatg	gaa gaacgggaag aggt			24
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40005				
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	escription of Artificial	sequence:	rck Primer	
s	equence			
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				2.4
cryggga	ggg tcaaagaagg agct			24

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                                                                    25
aaggctgggc ctaacccagt ctcat
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                                                                    25
gtccctgcag gagaagccag tgtac
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ctgggcaaat cetcaettge ttgtet	20
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<211> 26 <212> DNA	
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varav interfer bequeitee	
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	20
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<211> 10	
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<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence: SAGE library	
tag sequence	
4400 105	
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ayootytoyo	10
<210> 186	
<211> 79	
<212> PRT	
<213> Homo sapiens	

<400> 186

Cys Tyr His Gly Asn Gly Glu Asn Tyr Arg Gly Thr Ala Ser Thr Thr

1 5 10 15

Glu Ser Gly Ala Pro Cys Gln Arg Trp Asp Ser Gln Thr Pro His Arg
20 25 30

His Ser Lys Tyr Thr Pro Glu Arg Tyr Pro Ala Lys Gly Leu Gly Glu 35 40 45

Asn Tyr Cys Arg Asn Pro Asp Gly Asp Glu Arg Pro Trp Cys Tyr Thr 50 55 60

Thr Asp Pro Arg Val Arg Trp Glu Tyr Cys Asp Ile Pro Arg Cys 65 70 75

<210> 187

<211> 81

<212> PRT

<213> Homo sapiens

<400> 187

Cys Tyr Ala Gly Asn Gly Glu Ser Tyr Arg Gly Thr Ala Ser Thr Thr

1 5 10 15

Lys Ser Gly Lys Pro Cys Gln Arg Trp Asp Ser Gln Thr Pro His Leu 20 25 30

His Arg Phe Thr Pro Glu Arg Phe Pro Glu Leu Gly Leu Glu His Asn
35 40 45

Tyr Cys Arg Asn Pro Asp Gly Asp Ser Glu Gly Pro Trp Cys Tyr Thr
50 55 60

Thr Asp Pro Asn Val Arg Trp Glu Tyr Cys Asp Ile Pro Gln Cys Glu 65 70 75 80

Ser

<210> 188

<211> 81

<212> PRT

<213> Homo sapiens

<400> 188

Arg Asp Cys Tyr Ala Gly Asn Gly Glu Ser Tyr Arg Gly Thr Ala Ser 1 5 10 15

Thr Thr Lys Ser Gly Lys Pro Cys Gln Arg Trp Asp Ser Gln Thr Pro
20 25 30

His Leu His Arg Phe Thr Pro Glu Arg Phe Pro Glu Leu Gly Leu Glu 35 40 45

His Asn Tyr Cys Arg Asn Pro Asp Gly Asp Ser Glu Gly Pro Trp Cys 50 60

Tyr Thr Thr Asp Pro Asn Val Arg Trp Glu Tyr Cys Asp Ile Pro Gln 65 70 75 80

Cys

<210> 189

<211> 75

<212> PRT

<213> Homo sapiens

<400> 189

Cys Phe Val Arg Leu Pro Asn Thr Lys Leu Pro Asp Phe Ser Pro Ile
1 5 10 15

Val Ile Ser Val Ala Ser Leu Glu Glu Cys Ala Gln Lys Cys Leu Asn 20 25 30

Ser Asn Cys Ser Cys Arg Ser Phe Thr Tyr Asn Asn Asp Thr Lys Gly 35 40 45

Cys Leu Leu Trp Ser Glu Ser Ser Leu Gly Asp Ala Arg Gln Leu Leu 50 55 60

Pro Ser Gly Gly Val Asp Tyr Tyr Glu Lys Ile
65 70 75

<210> 190

<211> 181

<212> PRT

<213> Homo sapiens

<400> 190

Met Gly Lys Leu Val Ser Arg Arg Ser Tyr Glu His Trp Glu Tyr Leu

Thr Ala Thr Leu Ile Ser Ile Gly Val Ser Met Phe Leu Leu Ser Ser 20 25 30

Gly Pro Glu Pro Arg Ser Ser Pro Ala Thr Thr Leu Ser Gly Leu Ile 35 40 45

Leu Leu Ala Gly Tyr Ile Ala Phe Asp Ser Phe Thr Ser Asn Trp Gln 50 55 60

Asp Ala Leu Phe Ala Tyr Lys Met Ser Ser Val Gln Met Met Phe Gly 65 70 75 80

Val Asn Phe Phe Ser Cys Leu Phe Thr Val Gly Ser Leu Leu Glu Gln 85 90 95

Gly Ala Leu Leu Glu Gly Thr Arg Phe Met Gly Arg His Ser Glu Phe
100 105 110

Ala Ala His Ala Leu Leu Leu Ser Ile Cys Ser Ala Cys Gly Gln Leu 115 120 125

Phe Ile Phe Tyr Thr Ile Gly Gln Phe Gly Ala Ala Val Phe Thr Ile 130 135 140

Ile Met Thr Leu Arg Gln Ala Phe Ala Ile Leu Leu Ser Cys Leu Leu 145 150 155 160

Tyr Gly His Thr Val Thr Val Val Gly Gly Leu Gly Val Ala Val Val
165 170 175

Phe Ala Ala Leu Leu 180

<210> 191

<211> 68

<212> PRT

<213> Homo sapiens

<400> 191

Trp Tyr Ser Ser Pro Pro Leu Tyr Val Tyr Trp Phe Arg Asp Gly Glu

1 5 10 15

Ile Pro Tyr Tyr Ala Glu Val Val Ala Thr Asn Asn Pro Asp Arg Arg 20 25 30

Val Lys Pro Glu Thr Gln Gly Arg Phe Arg Leu Leu Gly Asp Val Gln 35 40 45

Lys Lys Asn Cys Ser Leu Ser Ile Gly Asp Ala Arg Met Glu Asp Thr 50 55 60

Gly Ser Tyr Phe 65